EXTENDING THE SEARCH FOR SYMMETRIES IN THE GENETIC CODE

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We report on the search for symmetries in the genetic code involving the medium rank simple Lie algebras $B_6 = \mathfrak{so}(13)$ and $D_7 = \mathfrak{so}(14)$, in the context of the algebraic approach originally proposed by one of the present authors, which aims at explaining the degeneracies encountered in the genetic code as the result of a sequence of symmetry breakings that have occurred during its evolution.

Keywords: Lie algebras; representation theory; genetic code.

1. Introduction

The genetic code constitutes a dictionary for protein synthesis that to each codon (triplet of nucleic bases in DNA or RNA) associates one of the twenty fundamental aminoacids or the termination signal. It summarizes, in condensed form, the rules governing the complex process of translation that occurs in the ribosome, involving all three forms of RNA (mRNA, tRNA and rRNA) together with various proteins and other auxiliary molecules. The degeneracy of this code, resulting simply from the fact that the number of aminoacids (20) is much smaller than that of codons (64), immediately suggests an analysis based on invariance principles, that is, on the concept of symmetry. The main idea is to construct a codon space, i.e. a 64-dimensional vector space carrying an irreducible representation of a group $G$. The irreducibility of this representation is lost, step by step, by imposing that in each step the symmetry is lowered to a maximal subgroup of $G$, which must generate precisely twenty-one subspaces

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to allocate the aminoacids and the termination signal. This process of spontaneous symmetry breaking has been used before, in various fields of science. The crucial point in such a search for symmetries is to find the ancestor group \( G \) and the possible chains of subgroups.

The first search for symmetries in the genetic code along this line of reasoning, as outlined in Ref. 1 (see also Ref. 2, and Refs. 3 and 4 for comments), has been carried out within the class of ordinary compact Lie groups \( G \) or, what amounts to the same thing, of semisimple Lie algebras \( \mathfrak{g} \), based on Cartan’s classification of semisimple Lie algebras, Dynkin’s classification of their maximal semisimple subalgebras\(^5,6\) and the tables of branching rules of McKay and Patera.\(^7\) This search has shown that, on the one hand, there is no simple Lie algebra that can generate the standard genetic code directly in its present form. On the other hand, it turned out that the standard genetic code can be obtained from the symplectic algebra \( \mathfrak{sp}(6) \) if, in the last step, the process of symmetry breaking is allowed to be gently interrupted, or frozen. This means that a few of the multiplets (subspaces) that have resulted from previous steps of the process do not participate in the symmetry reduction implied by the last step.

Two restrictions have been imposed on the search reported in Ref. 1. The first refers to the last phase of the process, after the ancestor algebra has already been broken to a direct sum of \( \mathfrak{su}(2) \)-subalgebras. In this last phase, further steps consist in breaking one or several of these \( \mathfrak{su}(2) \)-subalgebras completely, which in the language of atomic or nuclear physics can be implemented by introducing an appropriate generator \( L_z \) into the Hamiltonian. Another possibility that was also considered is to introduce instead its square \( L_z^2 \), which leads to a softer form of symmetry breaking. However, the possibility of performing both of these breakings sequentially, with freezing applied to the second step only, was not contemplated. The second restriction was the exclusion of “diagonal breaking”, which in the case of \( \mathfrak{su}(2) \)-subalgebras amounts to a Clebsch–Gordan summation law for quantum numbers.

In Ref. 8, this search has been enlarged as a consequence of a new geometric interpretation of the operator \( L_z^2 \) (rather than \( L_z \)); it implements symmetry breaking of the compact Lie group \( SU(2) \) to its maximal non-connected subgroup \( O(2) \) (rather than its maximal connected subgroup \( SO(2) \)). This point of view makes it perfectly acceptable to allow a symmetry breaking from \( SU(2) \) to \( O(2) \) followed by one from \( O(2) \) to \( SO(2) \). As a result, there appeared a new chain based on the exceptional group \( G_2 \). Soon after, a second chain based on \( G_2 \) was discovered. The main problem with both of these \( G_2 \) models, which will appear again in the analysis of the medium rank algebras \( \mathfrak{so}(13) \) and \( \mathfrak{so}(14) \) to be reported in this paper, is the large number of multiplets (subspaces) which must be frozen, in contrast to biological information that seems to speak in favor of a small amount of freezing.

A complete review of the full process has been given in Ref. 9, eliminating the restrictions pointed out above for the low rank algebras. Here, we extend this analysis to the medium rank algebras \( \mathfrak{so}(13) \) and \( \mathfrak{so}(14) \). In this sense, the present paper
is a companion of Ref. 9. Of course, we were aware from the very beginning that the analysis of hundreds of chains, required to handle this case, is a formidable task. However, when attempting to fully settle the question of the existence of symmetries behind a fundamental biological structure such as that underlying the organization of the genetic code, we cannot afford to discard any reasonable possibilities \textit{a priori}.

Another aspect that deserves to be commented is that, as already proposed in Ref. 1, a similar analysis for Lie superalgebras has already been completed,\textsuperscript{10,11} whereas the same investigation in the context of finite groups is under way and has so far given rise to partial results.\textsuperscript{12}

This article, which has grown out of previous work by Y.M.M. Hornos (unpublished) and the authors,\textsuperscript{13,14} is organized as follows. Section 2 is dedicated to exposing the general strategy adopted in the search that is reported in the following three sections. Given the fact that branching from the medium rank simple Lie algebras $B_6 = \mathfrak{so}(13)$ and $D_7 = \mathfrak{so}(14)$ leads to thousands of chains that are impossible to analyze case by case, our strategy is to first formulate a set of rules that allows one to discard the great majority of these chains without detailed analysis, thus reducing the number of "surviving" chains to manageable proportions. Sections 3–5 report the details of this search, organized in three phases and in a tree-like structure, where each branch is followed up to the point where it can either be discarded or else be shown to produce the pattern of degeneracies observed in the genetic code. In Section 6, we summarize the results, and in Section 7, we comment on conclusions that can be drawn from our work.

Given the highly technical nature of Sections 3–5, in which the material has been organized into enormous lists, the reader is strongly advised to concentrate first on Section 2 and then on Section 6, consulting Sections 3–5 according to necessity or to his/her personal preferences and interests, in an encyclopaedic manner.

\section{General Strategy}

The mathematical procedure for analyzing the degeneracies of the genetic code is based on repeated symmetry reduction to maximal subalgebras, leading to descending chains of subalgebras, each of which is maximal in the previous one. For this purpose, it is sufficient to consider only semisimple subalgebras, because a possible non-trivial center does not contribute to dimensions or branching rules. Therefore, the explicit construction of all possible chains requires, as a pre-requisite, a complete classification of the maximal semisimple subalgebras of semisimple Lie algebras — a problem solved long ago by Dynkin.\textsuperscript{5,6}

An important first step in Dynkin’s classification is to realize that the general problem can be reduced to that of classifying the maximal semisimple subalgebras of simple Lie algebras, namely by means of a theorem (Ref. 5, Theorem 15.1) stating that the maximal semisimple subalgebras $\mathfrak{g}'$ of a semisimple Lie algebra $\mathfrak{g}$ are of two types:
Correspondingly, the chains can be classified into simple chains, i.e. chains that do not involve breaking to a subalgebra of diagonal type at any point, and diagonal chains, i.e. chains that do so at some point.

The task of classifying the maximal semisimple subalgebras of simple Lie algebras is much more difficult and forms the core of Dynkin’s work. The results for the simple Lie algebras of interest here are assembled in Table 1; they can also be read off from the tables of McKay and Patera\textsuperscript{7} that are themselves, in this respect, derived from Dynkin’s work and have been verified independently by one of the present authors.\textsuperscript{14}

<table>
<thead>
<tr>
<th>Cartan label of simple Lie algebra</th>
<th>Cartan label of maximal semisimple subalgebra</th>
</tr>
</thead>
<tbody>
<tr>
<td>(D_7)</td>
<td>(A_6, B_6, C_3, C_2, G_2, A_3 \oplus D_4, A_1 \oplus B_5, C_2 \oplus B_4, B_3 \oplus B_5, A_1 \oplus A_1 \oplus D_5)</td>
</tr>
<tr>
<td>(A_6)</td>
<td>(A_5, B_3, A_1 \oplus A_4, A_2 \oplus A_3)</td>
</tr>
<tr>
<td>(B_6)</td>
<td>(D_6, A_1, A_1 \oplus D_5, C_2 \oplus D_4, A_3 \oplus B_5, A_1 \oplus A_1 \oplus B_4)</td>
</tr>
<tr>
<td>(D_6)</td>
<td>(A_5, B_5, A_3 \oplus A_3, A_1 \oplus B_4, C_2 \oplus B_3, A_1 \oplus C_3, A_1 \oplus A_1 \oplus A_1 \oplus D_4)</td>
</tr>
<tr>
<td>(A_5)</td>
<td>(A_4, A_3, C_3, A_2, A_1 \oplus A_3, A_2 \oplus A_2, A_1 \oplus A_2)</td>
</tr>
<tr>
<td>(B_5)</td>
<td>(A_4, B_4, C_2, A_1 \oplus B_3, C_2 \oplus C_2, A_1 \oplus A_1 \oplus A_3)</td>
</tr>
<tr>
<td>(D_5)</td>
<td>(A_4, C_2, A_1 \oplus A_2)</td>
</tr>
<tr>
<td>(A_4)</td>
<td>(D_4, A_1, A_1 \oplus A_3, A_1 \oplus A_1, A_1 \oplus A_1 \oplus C_2)</td>
</tr>
<tr>
<td>(B_4)</td>
<td>(B_4, A_2, A_1 \oplus C_2, A_1 \oplus A_1 \oplus A_1 \oplus A_1)</td>
</tr>
<tr>
<td>(A_3)</td>
<td>(A_2, C_2, A_1 \oplus A_1)</td>
</tr>
<tr>
<td>(B_3)</td>
<td>(G_2, A_3, A_1 \oplus A_1 \oplus A_1)</td>
</tr>
<tr>
<td>(C_3)</td>
<td>(A_2, A_1, A_1 \oplus C_2, A_1 \oplus A_1)</td>
</tr>
<tr>
<td>(A_2)</td>
<td>(A_1^{(1)}, A_1^{(2)})</td>
</tr>
<tr>
<td>(C_2)</td>
<td>(A_1, A_1 \oplus A_1)</td>
</tr>
<tr>
<td>(G_2)</td>
<td>(A_2, A_1, A_1 \oplus A_1)</td>
</tr>
</tbody>
</table>
Here and in what follows, the subalgebras are ordered according to the following criteria: first by increasing number of simple summands, then by decreasing total rank and finally by decreasing rank for the simple summand of highest rank.

The general strategy for analyzing the chains that result from repeated symmetry reduction to maximal semisimple subalgebras consists in proceeding along each chain one step at a time and analyzing, after each step, whether the resulting pattern of degeneracies is still compatible with that of the genetic code, as shown in Table 2. If not, the chain is non-surviving and may be discarded without further analysis. Otherwise, the chain is surviving (up to the stage considered), which means that we must proceed to analyze the possible next steps of the symmetry breaking.

To organize the multitude of chains that have to be analyzed, we have found it convenient to first of all divide the whole process into three distinct phases (rather than two, as stated in Ref. 9):

**Phase 1:** Breaking of the primordial symmetry to $\mathfrak{su}(2)$-symmetries.

During the first phase, symmetry breaking proceeds through chains of maximal semisimple subalgebras, either by simple breaking or by diagonal breaking of pairs of simple subalgebras different from $\mathfrak{su}(2)$. Every such chain will necessarily terminate in a direct sum of $p$ copies of $\mathfrak{su}(2)$, where $p$ may range from 1 up to the rank of the original simple Lie algebra.

**Phase 2:** Diagonal breaking of the $\mathfrak{su}(2)$-symmetries.

The second phase consists in subjecting the chains that have survived up to the end of phase 1 to diagonal breaking of the $\mathfrak{su}(2)$-summands. When there are $p$ such summands, there will a priori be $\binom{p}{2}$ possibilities of diagonal breaking, but this number can be substantially reduced when the previously obtained distribution of multiplets is invariant under certain permutations between the $\mathfrak{su}(2)$-summands.

**Phase 3:** Breaking of the $\mathfrak{su}(2)$-symmetries.

The third phase consists in subjecting the chains that have survived up to the end of phase 2 to breaking of (some of) the $\mathfrak{su}(2)$-subalgebras, either to $\mathfrak{o}(2)$ or to $\mathfrak{so}(2)$, including the possibility of freezing in the last step, as explained in Ref. 9.

**Table 2.** Dimensions and multiplicities in the standard genetic code.

<table>
<thead>
<tr>
<th>Dimension of multiplet</th>
<th>Number of multiplets</th>
<th>Amino acids</th>
</tr>
</thead>
<tbody>
<tr>
<td>6</td>
<td>3</td>
<td>Arg, Leu, Ser</td>
</tr>
<tr>
<td>4</td>
<td>5</td>
<td>Ala, Gly, Pro, Thr, Val</td>
</tr>
<tr>
<td>3</td>
<td>2</td>
<td>Ile, TERM</td>
</tr>
<tr>
<td>2</td>
<td>9</td>
<td>Asn, Asp, Cys, Gln, Glu, His, Lys, Phe, Tyr</td>
</tr>
<tr>
<td>1</td>
<td>2</td>
<td>Met, Trp</td>
</tr>
</tbody>
</table>
Next, it is useful to assemble the chains into families characterized by the same final algebra and the same distribution of multiplets under it. This reflects the fact that if

\[ g \supset g_1 \supset g_2 \supset \cdots \supset g_r \supset h \]

and

\[ g \supset g'_1 \supset g'_2 \supset \cdots \supset g'_r \supset h' \]

are two descending chains of subalgebras starting out from the same initial algebra \( g \) and ending in final subalgebras \( h \) and \( h' \) of \( g \) that are conjugate in \( g \), then the distributions of multiplets with respect to \( h \) and to \( h' \) obtained from any given representation of \( g \) along the two chains are identical — except when the last step of the symmetry reduction (from \( g_r \) to \( h \) and/or from \( g'_r \) to \( h' \)) involves freezing.\(^9\)

Of course, it is then sufficient to analyze the further fate of only one chain in each such family.

Our next task is to specify the criteria used to identify non-surviving chains. To this end, we observe first of all that there are a few simple rules that allow one to decide whether the symmetry breaking must be stopped at the stage considered or whether it must proceed at least one stage further. As explained in Ref. 9, the process stops whenever we encounter

- more than 21 multiplets,
- more than 2 singlets,
- more than 4 odd-dimensional multiplets,
- not enough multiplets of dimension \( \geq 6 \) or \( \geq 4 \),

whereas it must be continued whenever we encounter

- less than 21 multiplets,
- multiplets of dimension \( \geq 7 \),
- more than 3 multiplets of dimension 6,
- multiplets of dimension 5.

In the limiting case of exactly 21 multiplets, the chain must either reproduce precisely the degeneracies of the genetic code or else must be discarded.

The simplest of these conditions is certainly the one referring to the total number of multiplets. When it exceeds 21, this can be taken as an indication that we should discard the chain in question, return to the previous stage and look for other possibilities to proceed from there. In the following, we shall therefore apply the classification of chains as surviving or non-surviving only when the total number of multiplets does not exceed 21: all others could be considered as trivially non-surviving. There are then various properties that allow one to identify non-surviving chains:

- Total pairing: all multiplets occur in (identical or complex conjugate) pairs.
- Singlet excess: more than 2 singlets.
Odd multiplet excess: more than 4 odd-dimensional multiplets.

Large multiplet defect: not enough multiplets of dimension \( \geq 6 \) or \( \geq 4 \).

These criteria for exclusion of chains are “hereditary” in the sense of being based on features, not shared by the distribution of multiplets found in the genetic code, that cannot be removed by any further symmetry breaking and therefore allow one to discard the chain together with all of its possible descendants. Apart from these, there exist further criteria for the exclusion of chains which are only “partially hereditary” in the sense of allowing one to discard a certain chain and some but not all of its descendants; this is the case for a set of rules regarding the possibility of producing the 3 sextets and 2 triplets of the genetic code and collectively referred to as the sextet/triplet generating rules.

To explain the content of these rules, which apply to the third phase of the symmetry breaking process, assume that we are given a surviving chain (where “surviving” now means that it has survived up to the end of the first and/or second phase): it will necessarily end in the direct sum of, say, \( p \) copies of \( \mathfrak{su}(2) \) where, for chains starting out from the codon representations of \( B_6 = \mathfrak{so}(13) \) or \( D_7 = \mathfrak{so}(14) \), \( p \) may range from 1 up to 6. Each of the multiplets appearing in the final distribution of multiplets for any such chain is characterized by its highest weight, which can be written in the form \((k_1, \ldots, k_p)\) with non-negative integers \( k_1, \ldots, k_p \); its dimension is \((k_1 + 1) \cdots (k_p + 1)\). Suppose now that we perform a symmetry reduction by breaking the \( j \)-th \( \mathfrak{su}(2) \), that is, from \( \mathfrak{su}_1(2) \oplus \cdots \oplus \mathfrak{su}_p(2) \) either to \( \mathfrak{su}_1(2) \oplus \cdots \oplus \mathfrak{so}_j(2) \oplus \cdots \oplus \mathfrak{su}_p(2) \) or to \( \mathfrak{su}_1(2) \oplus \cdots \oplus \mathfrak{so}_j(2) \oplus \cdots \oplus \mathfrak{su}_p(2) \). In the first case, the multiplet \((k_1, \ldots, k_p)\) will split into \( k_j + 1 \) multiplets of dimension \((k_1 + 1) \cdots (k_{j-1} + 1) \cdot (k_{j+1} + 1) \cdots (k_p + 1)\), whereas in the second case, it will split into:

- \( \frac{1}{2}(k_j + 1) \) multiplets of dimension \( 2(k_1 + 1) \cdots (k_{j-1} + 1) \cdot (k_{j+1} + 1) \cdots (k_p + 1) \), if \( k_j \) is odd, and
- \( \frac{1}{2}k_j \) multiplets of dimension \( 2(k_1 + 1) \cdots (k_{j-1} + 1) \cdot (k_{j+1} + 1) \cdots (k_p + 1) \) plus 1 multiplet of dimension \((k_1 + 1) \cdots (k_{j-1} + 1) \cdot (k_{j+1} + 1) \cdots (k_p + 1)\), if \( k_j \) is even.

Repeating this process, we see that any breaking whatsoever of the multiplet \((k_1, \ldots, k_p)\) will lead to a distribution of sub-multiplets whose dimensions are products of certain of the numbers \( k_j + 1 \), corresponding to those \( \mathfrak{so}(2) \)'s that have remained unbroken, and when breaking to \( \mathfrak{so}(2) \) is involved, some power of 2. In particular, sub-multiplets whose dimension is a multiple of 3 can only appear if at least one of the numbers \( k_j + 1 \) is a multiple of 3 (and the corresponding \( \mathfrak{su}_j(2) \) is unbroken); moreover, sextets/triplets can only appear if at least one of them is actually equal to 6 or 3/3. Finally, observe that if any one of the integers \( k_j \) is greater than 5, then the corresponding \( \mathfrak{su}_j(2) \) must be broken, since otherwise there would remain multiplets of dimension \( \geq 7 \). Thus we arrive at
Rule 1: During the last phase of the symmetry breaking process, sub-multiplets whose dimension is a multiple of 3 can only appear as the result of breaking multiplets \((k_1, \ldots, k_p)\) whose dimension is a multiple of 3, that is, for which at least one of the integers \(k_1, \ldots, k_p\) equals 2 \(\mod\) 3. More specifically, the triplets come from multiplets \((k_1, \ldots, k_p)\) where at least one of the integers \(k_1, \ldots, k_p\) equals 2, whereas the sextets come from multiplets \((k_1, \ldots, k_p)\) where at least one of the integers \(k_1, \ldots, k_p\) equals 2 or 5.

Rule 2: During the last phase of the symmetry breaking process, breaking a multiplet whose dimension is a multiple of 3 results in a collection of sub-multiplets whose dimension is a multiple of 3 either for all or for none of the sub-multiplets within the collection.

Based on these rules, we can make a list of all multiplets \((k_1, \ldots, k_p)\) appearing in the final distribution of multiplets for one of the surviving chains for \(B_6 = \mathfrak{so}(13)\) or \(D_7 = \mathfrak{so}(14)\) whose dimension is a multiple of 3 and that after further breaking can produce sextets and/or triplets. This information is collected in Table 3, except that we omit breakings that do not produce any sub-multiplets whose dimension is a multiple of 3 (1 sextet into 6 singlets or 3 doublets, rather than 2 triplets, for example) or that do not succeed in eliminating all sub-multiplets of dimension > 6 (1 multiplet of dimension 24 into 2 multiplets of dimension 12, rather than 4 sextets or 8 triplets, for example). Without loss of generality, the integer that, according to

<table>
<thead>
<tr>
<th>Multiplet ((k_1, \ldots, k_p))</th>
<th>Dimension</th>
<th>Possible breakings into sextets/triplets</th>
</tr>
</thead>
<tbody>
<tr>
<td>((5, 0, \ldots))</td>
<td>6</td>
<td>1 sextet (unbroken)</td>
</tr>
<tr>
<td>((2, 0, \ldots))</td>
<td>3</td>
<td>1 triplet (unbroken)</td>
</tr>
<tr>
<td>((5, 1, 0, \ldots))</td>
<td>12</td>
<td>2 sextets</td>
</tr>
<tr>
<td>((2, 6, 0, \ldots))</td>
<td>21</td>
<td>3 sextets and 1 triplet</td>
</tr>
<tr>
<td></td>
<td></td>
<td>7 triplets</td>
</tr>
<tr>
<td>((2, 5, 0, \ldots))</td>
<td>18</td>
<td>3 sextets</td>
</tr>
<tr>
<td></td>
<td></td>
<td>6 triplets</td>
</tr>
<tr>
<td>((2, 4, 0, \ldots))</td>
<td>15</td>
<td>2 sextets and 1 triplet</td>
</tr>
<tr>
<td></td>
<td></td>
<td>5 triplets</td>
</tr>
<tr>
<td>((2, 3, 0, \ldots))</td>
<td>12</td>
<td>2 sextets</td>
</tr>
<tr>
<td></td>
<td></td>
<td>4 triplets</td>
</tr>
<tr>
<td>((2, 2, 0, \ldots))</td>
<td>9</td>
<td>1 sextet and 1 triplet</td>
</tr>
<tr>
<td></td>
<td></td>
<td>3 triplets</td>
</tr>
<tr>
<td>((2, 1, 0, \ldots))</td>
<td>6</td>
<td>1 sextet (unbroken)</td>
</tr>
<tr>
<td></td>
<td></td>
<td>2 triplets</td>
</tr>
<tr>
<td>((2, 1, 3, 0, \ldots))</td>
<td>24</td>
<td>4 sextets</td>
</tr>
<tr>
<td></td>
<td></td>
<td>8 triplets</td>
</tr>
<tr>
<td>((2, 1, 2, 0, \ldots))</td>
<td>18</td>
<td>3 sextets</td>
</tr>
<tr>
<td></td>
<td></td>
<td>6 triplets</td>
</tr>
<tr>
<td>((2, 1, 1, 0, \ldots))</td>
<td>12</td>
<td>2 sextets</td>
</tr>
<tr>
<td></td>
<td></td>
<td>4 triplets</td>
</tr>
</tbody>
</table>
rule 1, must be equal to 2 or 5 is assumed to be \( k_1 \), and when we write \((k_1, 0, \ldots), (k_1, k_2, 0, \ldots), (k_1, k_2, k_3, 0, \ldots), \) etc, the dots indicate that the remaining \( k \)'s are meant to be zero.

The simplest consequence of these sextet/triplet generating rules is of course the observation, already formalized in Ref. 9, that a chain will be non-surviving when there are not enough multiplets whose dimension is a multiple of 3: adding up the dimensions of all these multiplets gives a number called \( d_3 \) in Ref. 9 that must be at least 24 in order for the chain to be surviving.

### 3. First Phase of Symmetry Breaking

In what follows, we list the chains that result from carrying out the symmetry breaking up to the end of the first phase. Except when explicitly stated otherwise, the total number of multiplets obtained after the last step is always less than 21, so that according to the criteria stated above, the procedure must be continued; in particular no freezing is allowed in this phase. We also present the relevant information on the distribution of multiplets obtained after the last step that allows to discard the non-surviving chains.

\[ B_6 = so(13) \]

1. \( B_6 \supset D_6 \):

1.1. \( B_6 \supset D_6 \supset A_5 \): continuing the symmetry breaking process, we obtain the following chains, all of which can be excluded:
   1.1.1. \( B_6 \supset D_6 \supset A_5 \supset A_4 \): 4 quintets and 4 singlets, as well as total pairing.
   1.1.2. \( B_6 \supset D_6 \supset A_5 \supset A_3 \): total pairing.
   1.1.3. \( B_6 \supset D_6 \supset A_5 \supset C_3 \): 4 singlets.
   1.1.4. \( B_6 \supset D_6 \supset A_5 \supset A_2 \): total pairing.
   1.1.5. \( B_6 \supset D_6 \supset A_5 \supset A_1 \supset A_3 \): 4 singlets.
   1.1.6. \( B_6 \supset D_6 \supset A_5 \supset A_2 \supset A_3 \): 4 nonets, 8 triplets and 4 singlets.
   1.1.7. \( B_6 \supset D_6 \supset A_5 \supset A_1 \supset A_2 \): continuing the symmetry breaking process, we obtain the following chains:
      1.1.7.1. \( B_6 \supset D_6 \supset A_5 \supset A_1 \supset A_2 \) \( \supset A_1 \) \( \supset A_1 \): this chain exhibits precisely 21 multiplets: 3 sextets, 5 quartets, 4 triplets, 5 doublets and 4 singlets.
      1.1.7.2. \( B_6 \supset D_6 \supset A_5 \supset A_1 \supset A_2 \) \( \supset A_1 \) \( \supset A_1 \): 2 nonets, 2 quintets and 4 singlets.

1.2. \( B_6 \supset D_6 \supset B_3 \): total pairing.

1.3. \( B_6 \supset D_6 \supset A_3 \oplus A_3 \): total pairing.

1.4. \( B_6 \supset D_6 \supset A_1 \oplus B_4 \): total pairing.

1.5. \( B_6 \supset D_6 \supset C_2 \oplus B_3 \): total pairing.

1.6. \( B_6 \supset D_6 \supset A_1 \oplus C_3 \): continuing the symmetry breaking process, we obtain the following chains:
   1.6.1. \( B_6 \supset D_6 \supset A_1 \oplus C_3 \supset A_1 \oplus A_2 \):
      distribution of multiplets identical to that from the chain 1.1.7:
      \( B_6 \supset D_6 \supset A_5 \supset A_1 \oplus A_2 \).
we either perform a diagonal breaking of the distribution of multiplets identical to that from the chain 1.6.5: remaining.

4.3.2. 4.3.1.

3.1. 3.1.3.

3.1.5. 3.1.4.

3.1.3. 3.1.2.

3.1.2. 3.1.1.

we either perform a diagonal breaking of the distribution of multiplets identical to that from the chain 1.6.5: remaining.

4.4.2. 4.4.1.

4.4.1. 4.4.2.

4.3.1. 4.3.2.

4.3.2. 4.3.1.

3.1.5. 3.1.4.

3.1.4. 3.1.3.

3.1.3. 3.1.2.

3.1.2. 3.1.1.

we either perform a diagonal breaking of the distribution of multiplets identical to that from the chain 1.6.5: remaining.

4.4.2. 4.4.1.

4.4.1. 4.4.2.

4.3.1. 4.3.2.

4.3.2. 4.3.1.

3.1.5. 3.1.4.

3.1.4. 3.1.3.

3.1.3. 3.1.2.

3.1.2. 3.1.1.

we either perform a diagonal breaking of the distribution of multiplets identical to that from the chain 1.6.5: remaining.
5. $B_6 \supset A_3 \oplus B_3$: total pairing.
6. $B_6 \supset A_1 \oplus A_1 \oplus B_3$: continuing the symmetry breaking process, we obtain the following chains:
   6.1. $B_6 \supset A_1 \oplus A_1 \oplus B_4 \supset A_1 \oplus A_1 \oplus D_4$:
       distribution of multiplets identical to that from the chain 1.7:
       $B_6 \supset D_6 \supset A_1 \oplus A_1 \oplus D_4$.
   6.2. $B_6 \supset A_1 \oplus A_1 \oplus B_4 \supset A_1 \oplus A_1 \oplus A_1$: surviving.
   6.3. $B_6 \supset A_1 \oplus A_1 \oplus B_4 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_3$: total pairing.
   6.4. $B_6 \supset A_1 \oplus A_1 \oplus B_4 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1$: surviving.
   6.5. $B_6 \supset A_1 \oplus A_1 \oplus B_4 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus C_2$:
       distribution of multiplets identical to that from the chain 4.4:
       $B_6 \supset C_2 \oplus D_4 \supset C_2 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1$.

$D_7 = \mathfrak{so}(14)$

As a preliminary remark, we observe that the maximal subalgebras $B_6$, $C_3$, $C_2$ and $G_2$ of $D_7$ can be disregarded because the codon representations of $D_7$ remain irreducible under restriction to these subalgebras, so that these cases are covered by our previous analysis here ($B_6$) and in Ref. 9 ($C_3$, $C_2$ and $G_2$).

1. $D_7 \supset A_6$
   1.1. $D_7 \supset A_6 \supset A_5$: distribution of multiplets identical to that from the $B_6$-chain 1.1:
       $B_6 \supset D_6 \supset A_5$.
   1.2. $D_7 \supset A_6 \supset B_3$: continuing the symmetry breaking process, we obtain the following chains, all of which can be excluded:
      1.2.1. $D_7 \supset A_6 \supset B_3 \supset A_3$: total pairing.
      1.2.2. $D_7 \supset A_6 \supset B_3 \supset G_2$:
             1 multiplet of dimension 27, 3 septets and 2 singlets.
      1.2.3. $D_7 \supset A_6 \supset B_3 \supset A_1 \oplus A_1 \oplus A_1$: 2 nonets, 4 triplets and 2 singlets.
   1.3. $D_7 \supset A_6 \supset A_2 \oplus A_3$: continuing the symmetry breaking process, we obtain the following chains, all of which can be excluded:
      1.3.1. $D_7 \supset A_6 \supset A_2 \oplus A_3 \supset A_1^{(1)} \oplus A_3$: 4 singlets.
      1.3.2. $D_7 \supset A_6 \supset A_2 \oplus A_3 \supset A_1^{(2)} \oplus A_3$: any posterior breaking of the $A_3$-algebra
       leads to a distribution of multiplets identical with that created by posterior
       breaking of the $A_2$-algebra in one of the following three cases and thus contains
       far too many odd-dimensional multiplets.
      1.3.3. $D_7 \supset A_6 \supset A_2 \oplus A_3 \supset A_2 \oplus A_2$:
             4 nonets, 8 triplets and 4 singlets.
      1.3.4. $D_7 \supset A_6 \supset A_2 \oplus A_3 \supset A_2 \oplus C_2$:
             1 multiplet of dimension 15, 1 quintet, 3 triplets and 3 singlets.
      1.3.5. $D_7 \supset A_6 \supset A_2 \oplus A_3 \supset A_2 \oplus A_1 \oplus A_1$: 2 nonets, 4 triplets and 2 singlets.
   1.4. $D_7 \supset A_6 \supset A_1 \oplus A_4$: continuing the symmetry breaking process, we obtain the following chains, all of which can be excluded:
      1.4.1. $D_7 \supset A_6 \supset A_1 \oplus A_4 \supset A_1 \oplus A_3$: 4 singlets.
      1.4.2. $D_7 \supset A_6 \supset A_1 \oplus A_4 \supset A_1 \oplus C_2 \supset A_1 \oplus A_1$:
             2 septets, 2 quintets, 2 triplets and 2 singlets.
      1.4.3. $D_7 \supset A_6 \supset A_1 \oplus A_4 \supset A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1$: 4 triplets and 4 singlets.
      1.4.4. $D_7 \supset A_6 \supset A_1 \oplus A_4 \supset A_1 \oplus A_1 \oplus A_2$: 4 triplets and 4 singlets.
2. $D_7 \supset A_3 \oplus D_4$: in the next step of the symmetry breaking process, we break the $D_4$-subalgebra, observing that this must be done at some stage since otherwise the dimensions of all multiplets would remain multiples of 8; this leads to the following chains:

2.1. $D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus B_3$: total pairing.

2.2. $D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_2$: total pairing.

2.3. $D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2$: in the next step of the symmetry breaking process, we break the $A_3$-subalgebra, observing that this must be done at some stage since otherwise the dimensions of all multiplets would remain multiples of 4; this leads to the following chains:

2.3.1. $D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2 \supset A_2 \oplus A_1 \oplus C_2 \supset A_1^{(1)} \oplus A_1 \oplus A_1$: surviving.

2.3.2. $D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2 \supset A_2 \oplus A_1 \oplus C_2 \supset A_1^{(2)} \oplus A_1 \oplus A_1$: surviving.

2.3.3. $D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2 \supset A_2 \oplus A_1 \oplus C_2 \supset A_1^{(3)} \oplus A_1 \oplus A_1 \oplus A_1$: surviving.

2.3.4. $D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2 \supset A_2 \oplus A_1 \oplus C_2 \supset A_1^{(4)} \oplus A_1 \oplus A_1 \oplus A_1$: surviving.

2.4. $D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus A_1 \oplus A_1$: continuing the symmetry breaking process, we obtain the following chains:

2.4.1. $D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1$: continuing the symmetry breaking process, we obtain the following chains:

2.4.1.1. $D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1$: continuing the symmetry breaking process, we obtain the following chains:

2.4.1.2. $D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1$: surviving.

2.4.2. $D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1$: distribution of multiplets identical to that from the $B_6$-chain 4.4:

$B_6 \supset C_2 \supset D_4 \supset C_2 \oplus A_1 \oplus C_2$.

2.4.3. $D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1$: surviving.

3. $D_7 \supset A_1 \oplus B_5$: continuing the symmetry breaking process, we obtain the following chains:

3.1. $D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus D_5$: total pairing.
3.2. \( D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1 \): surviving.

3.3. \( D_7 \supset A_1 \oplus B_5 \oplus A_1 \oplus A_1 \oplus D_4 \): continuing the symmetry breaking process, we obtain the following chains:

3.3.1. \( D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus B_3 \): total pairing.

3.3.2. \( D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1 \oplus D_4 \supset A_1 \oplus A_1 \oplus A_2 \): total pairing.

3.3.3. \( D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1 \oplus D_4 \supset A_1 \oplus A_1 \oplus A_1 \oplus C_2 \):

distribution of multiplets identical to that from the chain 2.3.3:

\( D_7 \supset A_3 \oplus D_4 \oplus A_3 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus C_2 \).

3.3.4. \( D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1 \oplus D_4 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \):

distribution of multiplets identical to that from the chain 2.4.3:

\( D_7 \supset A_3 \oplus D_4 \oplus A_3 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \).

3.4. \( D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1 \oplus C_2 \): total pairing.

3.5. \( D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1 \oplus A_1 \oplus B_3 \): continuing the symmetry breaking process, we obtain the following chains:

3.5.1. \( D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \): total pairing.

3.5.2. \( D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus G_2 \): continuing the symmetry breaking process, we obtain the following chains:

3.5.2.1. \( D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1 \oplus A_1 \oplus B_3 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus G_2 \)

\( \supset A_1 \oplus A_1 \oplus A_1 \oplus A_2 \): total pairing.

3.5.2.2. \( D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1 \oplus A_1 \oplus B_3 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus G_2 \)

\( \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \): surviving.

3.5.2.3. \( D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus B_3 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus G_2 \)

\( \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \): surviving.

3.5.3. \( D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus B_3 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \):

distribution of multiplets identical to that from the chain 2.4.3:

\( D_7 \supset A_3 \oplus D_4 \oplus A_3 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \).

4. \( D_7 \supset C_2 \oplus B_4 \): in the next step of the symmetry breaking process, we break the \( B_4 \)-subalgebra, observing that this must be done at some stage since otherwise the dimensions of all multiplets would remain multiples of 16; this leads to the following chains:

4.1. \( D_7 \supset C_2 \oplus B_4 \oplus C_2 \oplus D_4 \): in the next step of the symmetry breaking process, we break the \( D_4 \)-subalgebra, observing that this must be done at some stage since otherwise the dimensions of all multiplets would remain multiples of 8; this leads to the following chains:

4.1.1. \( D_7 \supset C_2 \oplus B_4 \oplus C_2 \oplus D_4 \oplus C_2 \oplus B_3 \): total pairing.

4.1.2. \( D_7 \supset C_2 \oplus B_4 \oplus C_2 \oplus D_4 \oplus C_2 \oplus A_2 \): total pairing.

4.1.3. \( D_7 \supset C_2 \oplus B_4 \oplus C_2 \oplus D_4 \oplus C_2 \oplus A_1 \oplus C_2 \):

distribution of multiplets identical to that from the \( B_5 \)-chain 4.3:

\( B_4 \supset C_2 \oplus D_4 \oplus C_2 \oplus A_1 \oplus C_2 \).

4.1.4. \( D_7 \supset C_2 \oplus B_4 \oplus C_2 \oplus D_4 \oplus C_2 \oplus A_1 \oplus A_1 \oplus A_1 \):

distribution of multiplets identical to that from the \( B_5 \)-chain 4.4:

\( B_6 \supset C_2 \oplus D_4 \oplus C_2 \oplus A_1 \oplus A_1 \oplus A_1 \).

4.2. \( D_7 \supset C_2 \oplus B_4 \oplus C_2 \oplus A_1 \): continuing the symmetry breaking process, we obtain the following chains:

4.2.1. \( D_7 \supset C_2 \oplus B_4 \oplus C_2 \oplus A_1 \oplus A_1 \): surviving.

4.2.2. \( D_7 \supset C_2 \oplus B_4 \oplus C_2 \oplus A_1 \oplus A_1 \oplus A_1 \):

distribution of multiplets identical to that from the \( B_6 \)-chain 6.2:

\( B_6 \supset A_1 \oplus A_1 \oplus B_4 \oplus A_1 \oplus A_1 \oplus A_1 \).
4.3. $D_7 \supset C_2 \oplus B_4 \supset C_2 \oplus A_1 \oplus A_1$; total pairing.

4.4. $D_7 \supset C_2 \oplus B_4 \supset C_2 \oplus A_1 \oplus A_1$: continuing the symmetry breaking process, we obtain the following chains:

4.4.1. $D_7 \supset C_2 \oplus B_4 \supset C_2 \oplus A_1 \oplus A_1 \supset A_1 \oplus A_1 \oplus A_1$: surviving.

4.4.2. $D_7 \supset C_2 \oplus B_4 \supset C_2 \oplus A_1 \oplus A_1 \supset A_1 \oplus A_1 \oplus A_1$:

- distribution of multiplets identical to that from the $B_6$-chain 6.4:
  
  $B_6 \supset A_1 \oplus A_1 \oplus B_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1$.

4.5. $D_7 \supset C_2 \oplus B_4 \supset C_2 \oplus A_1 \oplus A_1 \oplus C_2$: continuing the symmetry breaking process, we either perform a diagonal breaking of the $C_2$-subalgebras and then break the resulting diagonal $C_2$-subalgebra, or else we break first one of the two $C_2$-subalgebras and then the other one; this leads to the following chains:

4.5.1. $D_7 \supset C_2 \oplus B_4 \supset C_2 \oplus A_1 \oplus A_1 \oplus C_2 \supset (C_2)_{14} \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1$:

- $A_1 \oplus A_1 \oplus A_1$; surviving.

4.5.2. $D_7 \supset C_2 \oplus B_4 \supset C_2 \oplus A_1 \oplus A_1 \oplus C_2 \supset (C_2)_{14} \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1$

- $A_1 \oplus A_1 \oplus A_1$; surviving.

4.5.3. $D_7 \supset C_2 \oplus B_4 \supset C_2 \oplus A_1 \oplus A_1 \oplus C_2 \supset C_2 \oplus A_1 \oplus A_1 \oplus A_1$

- $A_1 \oplus A_1 \oplus A_1$; surviving.

4.5.4. $D_7 \supset C_2 \oplus B_4 \supset C_2 \oplus A_1 \oplus A_1 \oplus C_2 \supset C_2 \oplus A_1 \oplus A_1 \oplus A_1$

- $A_1 \oplus A_1 \oplus A_1 \oplus A_1$:

- distribution of multiplets identical to that from the chain 2.4.2.1:

  $D_7 \supset A_3 \oplus B_4 \supset A_3 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus C_2 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1$

- $A_1 \oplus A_1 \oplus A_1 \oplus A_1$.

4.5.5. $D_7 \supset C_2 \oplus B_4 \supset C_2 \oplus A_1 \oplus A_1 \oplus C_2 \supset C_2 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1$

- $A_1 \oplus A_1 \oplus A_1 \oplus A_1$:

- distribution of multiplets identical to that from the chain 2.4.2.1:

  $D_7 \supset A_3 \oplus B_4 \supset A_3 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus C_2 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1$

- $A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1$.

4.5.6. $D_7 \supset C_2 \oplus B_4 \supset C_2 \oplus A_1 \oplus A_1 \oplus C_2 \supset C_2 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1$

- $A_1 \oplus A_1 \oplus A_1 \oplus A_1$:

- distribution of multiplets identical to that from the chain 2.4.2.2:

  $D_7 \supset A_3 \oplus B_4 \supset A_3 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus C_2 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1$

- $A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1$.

5. $D_7 \supset B_3 \oplus B_3$: continuing the symmetry breaking process, we obtain the following chains:

5.1. $D_7 \supset B_3 \oplus B_3 \supset B_3 \oplus A_3$: total pairing.

5.2. $D_7 \supset B_3 \oplus B_3 \supset B_3 \oplus G_2$: in the next step of the symmetry breaking process, we break the $B_3$-subalgebra, observing that this must be done at some stage since otherwise the dimensions of all multiplets would remain multiples of 8; this leads to the following chains:

5.2.1. $D_7 \supset B_3 \oplus B_3 \supset B_3 \oplus G_2 \supset A_3 \oplus G_2$: total pairing.

5.2.2. $D_7 \supset B_3 \oplus B_3 \supset B_3 \oplus G_2 \supset G_2 \oplus G_2$: continuing the symmetry breaking process, we either perform a diagonal breaking of the $G_2$-subalgebras or else we break first one of the two $G_2$-subalgebras and then the other one; this leads to the following chains:

5.2.2.1. $D_7 \supset B_3 \oplus B_3 \supset B_3 \oplus G_2 \supset G_2 \oplus G_2 \supset (G_2)_{12}$:

- 1 multiplet of dimension 27, 3 septets and 2 singlets.

5.2.2.2. $D_7 \supset B_3 \oplus B_3 \supset B_3 \oplus G_2 \supset G_2 \oplus G_2 \supset G_2 \oplus A_1 \oplus A_1 \oplus A_1$:

- surviving.
5.2.2.3. \( D_7 \supset B_3 \oplus B_3 \supset B_3 \oplus G_2 \supset G_2 \oplus G_2 \supset G_2 \oplus A_1 \supset A_1 \oplus A_1 \oplus A_1 \): surviving.

5.2.2.4. \( D_7 \supset B_3 \oplus B_3 \supset B_3 \oplus G_2 \supset G_2 \oplus G_2 \supset G_2 \oplus A_1 \oplus A_1 \supset A_1 \oplus A_1 \oplus A_1 \): surviving.

5.2.2.5. \( D_7 \supset B_3 \oplus B_3 \supset B_3 \oplus G_2 \supset G_2 \oplus G_2 \supset G_2 \oplus A_1 \oplus A_1 \supset A_1 \oplus A_1 \oplus A_1 \): surviving.

5.2.3. \( D_7 \supset B_3 \oplus B_3 \supset B_3 \oplus G_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus G_2 \):
distribution of multiplets identical to that from the chain 3.5.2:
\( D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1 \oplus A_1 \oplus B_3 \supset A_1 \oplus A_1 \oplus A_1 \oplus G_2 \).

5.3. \( D_7 \supset B_3 \oplus B_3 \supset B_3 \oplus A_1 \oplus A_1 \oplus A_1 \):
distribution of multiplets identical to that from the chain 3.5:
\( D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1 \oplus A_1 \oplus B_3 \).

5.4. \( D_7 \supset B_3 \oplus B_3 \supset (B_3)_12 \):
distribution of multiplets identical to that from the chain 1.2:
\( D_7 \supset A_6 \supset B_3 \).

6. \( D_7 \supset A_1 \oplus A_1 \oplus D_5 \): continuing the symmetry breaking process, we obtain the following chains:

6.1. \( D_7 \supset A_1 \oplus A_1 \oplus D_5 \supset A_1 \oplus A_1 \oplus A_1 \):
continuing the symmetry breaking process, we obtain the following chains:

6.1.1. \( D_7 \supset A_1 \oplus A_1 \oplus D_5 \supset A_1 \oplus A_1 \oplus A_1 \supset A_1 \oplus A_1 \oplus A_3 \supset A_1 \oplus A_1 \oplus A_2 \): total pairing.

6.1.2. \( D_7 \supset A_1 \oplus A_1 \oplus D_5 \supset A_1 \oplus A_1 \oplus A_4 \supset A_1 \oplus A_1 \oplus A_3 \supset A_1 \oplus A_1 \oplus A_2 \):
distribution of multiplets identical to that from the chain 4.5.2:
\( D_7 \supset C_2 \oplus B_3 \supset C_2 \oplus A_1 \oplus A_1 \oplus C_2 \supset (C_2)_14 \oplus A_1 \oplus A_1 \supset A_1 \oplus A_1 \oplus A_1 \).

6.1.2. \( D_7 \supset A_1 \oplus A_1 \oplus D_5 \supset A_1 \oplus A_1 \oplus A_4 \supset A_1 \oplus A_1 \oplus A_3 \supset A_1 \oplus A_1 \oplus A_1 \):
distribution of multiplets identical to that from the chain 4.5.1:
\( D_7 \supset C_2 \oplus B_4 \supset C_2 \oplus A_1 \oplus A_1 \oplus C_2 \supset (C_2)_14 \oplus A_1 \oplus A_1 \supset A_1 \oplus A_1 \oplus A_1 \).

6.1.3. \( D_7 \supset A_1 \oplus A_1 \oplus D_5 \supset A_1 \oplus A_1 \oplus A_4 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \):
distribution of multiplets identical to that from the chain 4.5.2:
\( D_7 \supset C_2 \oplus B_4 \supset C_2 \oplus A_1 \oplus A_1 \oplus C_2 \supset (C_2)_14 \oplus A_1 \oplus A_1 \supset A_1 \oplus A_1 \oplus A_1 \).

6.1.4. \( D_7 \supset A_1 \oplus A_1 \oplus D_5 \supset A_1 \oplus A_1 \oplus A_4 \supset A_1 \oplus A_1 \oplus A_2 \):
continuing the symmetry breaking process, we obtain the following chains:

6.1.4.1. \( D_7 \supset A_1 \oplus A_1 \oplus D_5 \supset A_1 \oplus A_1 \oplus A_4 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \):
non-surviving
(only 2 octets that by themselves cannot produce 3 sextets, and no other multiplets of dimension > 4).
6.1.4.2. \( D_7 \supset A_1 \otimes A_1 \otimes D_5 \supset A_1 \otimes A_1 \otimes A_4 \supset A_1 \otimes A_1 \otimes A_1 \otimes A_2 \)
\( \supset A_1 \otimes A_1 \otimes A_1 \otimes A_1^{(2)} \): surviving.

6.2. \( D_7 \supset A_1 \otimes A_1 \otimes D_5 \supset A_1 \otimes A_1 \otimes B_4 \): continuing the symmetry breaking process, we obtain the following chains:

6.2.1. \( D_7 \supset A_1 \otimes A_1 \otimes D_5 \supset A_1 \otimes A_1 \otimes B_4 \supset A_1 \otimes A_1 \otimes D_4 \): continuing the symmetry breaking process, we obtain the following chains:

6.2.1.1. \( D_7 \supset A_1 \otimes A_1 \otimes D_5 \supset A_1 \otimes A_1 \otimes B_4 \supset A_1 \otimes A_1 \otimes D_4 \supset A_1 \otimes A_1 \otimes B_3 \): total pairing.

6.2.1.2. \( D_7 \supset A_1 \otimes A_1 \otimes D_5 \supset A_1 \otimes A_1 \otimes B_4 \supset A_1 \otimes A_1 \otimes D_4 \supset A_1 \otimes A_1 \otimes A_2 \): total pairing.

6.2.1.3. \( D_7 \supset A_1 \otimes A_1 \otimes D_5 \supset A_1 \otimes A_1 \otimes B_4 \supset A_1 \otimes A_1 \otimes D_4 \)
\( \supset A_1 \otimes A_1 \otimes A_1 \otimes C_2 \supset A_1 \otimes A_1 \otimes A_1 \otimes A_1 \):

distribution of multiplets identical to that from the \( B_6 \)-chain 4.3.3:

\( B_6 \supset C_2 \otimes D_4 \supset C_2 \otimes A_1 \otimes C_2 \supset (A_1 \otimes A_1) \oplus A_1 \oplus A_1 \).

6.2.1.4. \( D_7 \supset A_1 \otimes A_1 \otimes D_5 \supset A_1 \otimes A_1 \otimes B_4 \supset A_1 \otimes A_1 \otimes D_4 \)
\( \supset A_1 \otimes A_1 \otimes A_1 \otimes C_2 \supset A_1 \otimes A_1 \otimes A_1 \otimes A_1 \):

distribution of multiplets identical to that from the \( B_6 \)-chain 4.3.5:

\( B_6 \supset C_2 \otimes D_4 \supset C_2 \otimes A_1 \otimes C_2 \supset A_1 \otimes A_1 \otimes A_1 \oplus A_1 \oplus A_1 \).

6.2.1.5. \( D_7 \supset A_1 \otimes A_1 \otimes D_5 \supset A_1 \otimes A_1 \otimes B_4 \supset A_1 \otimes A_1 \otimes D_4 \)
\( \supset A_1 \otimes A_1 \otimes A_1 \otimes A_1 \oplus A_1 \):

distribution of multiplets identical to that from the chain 2.4.3:

\( D_7 \supset A_3 \otimes D_4 \supset A_3 \otimes A_1 \oplus A_1 \oplus A_1 \oplus A_1 \)
\( \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \):

distribution of multiplets identical to that from the chain 4.2.2:

\( D_7 \supset C_2 \otimes B_4 \supset C_2 \otimes A_1 \oplus A_1 \oplus A_1 \).

6.2.2. \( D_7 \supset A_1 \otimes A_1 \otimes D_5 \supset A_1 \otimes A_1 \otimes B_4 \supset A_1 \otimes A_1 \oplus A_1 \):

distribution of multiplets identical to that from the chain 4.2.2:

\( D_7 \supset A_1 \otimes A_1 \otimes D_5 \supset A_1 \otimes A_1 \otimes B_4 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \).

6.2.3. \( D_7 \supset A_1 \otimes A_1 \otimes D_5 \supset A_1 \otimes A_1 \otimes B_4 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_3 \): total pairing.

6.2.4. \( D_7 \supset A_1 \otimes A_1 \otimes D_5 \supset A_1 \otimes A_1 \otimes B_4 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \):

distribution of multiplets identical to that from the chain 4.4.2:

\( D_7 \supset C_2 \otimes B_4 \supset C_2 \otimes A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \).

6.2.5. \( D_7 \supset A_1 \otimes A_1 \otimes D_5 \supset A_1 \otimes A_1 \otimes B_4 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus C_2 \):

distribution of multiplets identical to that from the chain 2.4.2:

\( D_7 \supset A_3 \otimes D_4 \supset A_3 \otimes A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \).

6.3. \( D_7 \supset A_1 \otimes A_1 \otimes D_5 \supset A_1 \otimes A_1 \otimes C_2 \): continuing the symmetry breaking process, we obtain the following chains:

6.3.1. \( D_7 \supset A_1 \otimes A_1 \otimes D_5 \supset A_1 \otimes A_1 \oplus C_2 \oplus A_1 \otimes A_1 \):

6.3.2. \( D_7 \supset A_1 \otimes A_1 \otimes D_5 \supset A_1 \otimes A_1 \oplus A_1 \otimes C_2 \):

6.4. \( D_7 \supset A_1 \otimes A_1 \otimes D_5 \supset A_1 \otimes A_1 \otimes B_3 \):

distribution of multiplets identical to that from the chain 3.5:

\( D_7 \supset A_1 \otimes B_5 \supset A_1 \otimes A_1 \otimes A_1 \oplus B_3 \).

6.5. \( D_7 \supset A_1 \otimes A_1 \otimes D_5 \supset A_1 \otimes A_1 \oplus C_2 \oplus C_2 \):

distribution of multiplets identical to that from the chain 4.5:

\( D_7 \supset C_2 \otimes B_4 \supset C_2 \otimes A_1 \oplus A_1 \oplus C_2 \).

6.6. \( D_7 \supset A_1 \otimes A_1 \otimes D_5 \supset A_1 \otimes A_1 \oplus A_1 \oplus A_1 \oplus A_3 \):

distribution of multiplets identical to that from the chain 2.4:

\( D_7 \supset A_3 \otimes D_4 \supset A_3 \otimes A_1 \oplus A_1 \oplus A_1 \oplus A_1 \).
4. Second Phase of Symmetry Breaking

In this section we list the chains that result from the surviving chains of phase one by applying diagonal breaking in all possible ways, except that chains obtained from each other by permutation of the $A_1$-summands are written only once. We also recall that if a chain is identified as non-surviving due to the sextet/triplet generating rules, it may still give rise to surviving descendants, obtained by further diagonal breaking.

$B_6 = \mathfrak{so}(13)$

- Chain 1.6.2:
  
  $B_6 \supset D_6 \supset A_1 \oplus C_3 \supset A_1 \oplus A_1$: non-surviving.

  This chain may be discarded due to the sextet/triplet generating rules in view of the fact that the two multiplets of dimension 18 that occur and that must of course be broken are incapable of generating the three sextets and two triplets in the genetic code: one of them has highest weight (1-8) and thus cannot produce any sextets or triplets at all while the other one has highest weight (2-5) and thus can only produce three sextets or six triplets or no sextets and triplets at all.

  1.6.2.1 $\cdots \supset (A_1)_{12}$: does not reproduce the genetic code.

- Chain 1.6.3:
  
  $B_6 \supset D_6 \supset A_1 \oplus C_3 \supset A_1 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1$: non-surviving.

  This chain may be discarded due to the sextet/triplet generating rules since it exhibits only one multiplet of dimension 12 and one sextet ($d_3 = 18$).

  1.6.3.1 $\cdots \supset (A_1)_{12} \oplus A_1$: total pairing.

  1.6.3.2 $\cdots \supset (A_1)_{13} \oplus A_1$: total pairing.

  1.6.3.3 $\cdots \supset (A_1)_{23} \oplus A_1$: surviving.

    1.6.3.3.1 $\cdots \supset (A_1)_{12}$: does not reproduce the genetic code.

- Chain 1.6.4:
  
  $B_6 \supset D_6 \supset A_1 \oplus C_3 \supset A_1 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1$: non-surviving.

  This chain may be discarded due to the sextet/triplet generating rules since it exhibits only three sextets ($d_3 = 18$).

  1.6.4.1 $\cdots \supset (A_1)_{12} \oplus A_1 \oplus A_1$: total pairing.

  1.6.4.2 $\cdots \supset (A_1)_{23} \oplus A_1 \oplus A_1$: surviving.

    1.6.4.2.1 $\cdots \supset (A_1)_{12} \oplus A_1$: total pairing.

    1.6.4.2.2 $\cdots \supset (A_1)_{13} \oplus A_1$: non-surviving.

    This chain may be discarded due to the sextet/triplet generating rules in view of the fact that the six sextets that occur and that may or may not be broken are incapable of generating the three sextets and two triplets in the genetic code: they are organized into three identical sextets with highest weight (2-1) plus another three identical sextets with highest weight (1-2) and thus cannot be broken so as to produce precisely two triplets.

    1.6.4.2.2.1 $\cdots \supset (A_1)_{12}$: does not reproduce the genetic code.

  1.6.4.2.3 $\cdots \supset (A_1)_{23} \oplus A_1$: total pairing.
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• Chain 1.6.5:

\[ B_0 \supset D_0 \supset A_1 \oplus C_3 \supset A_1 \oplus A_1 \oplus A_1: \text{non-surviving.} \]

This chain may be discarded due to the sextet/triplet generating rules in view of the fact that the two multiplets of dimension 18 that occur and that must of course be broken are incapable of generating the three sextets and two triplets in the genetic code: they have highest weights (2-1-2) and (1-2-2) and so each of them can only produce three sextets or two sextets and two triplets or six triplets or no sextets and triplets at all, which cannot be combined to yield three sextets and two triplets.

1.6.5.1 \( \cdots \supset (A_1)_{12} \oplus A_1: \text{total pairing.} \)

1.6.5.2 \( \cdots \supset (A_1)_{13} \oplus A_1: \) distribution of multiplets identical to that from the chain 1.6.3.3.

• Chain 1.7.3:

\[ B_0 \supset D_0 \supset A_1 \oplus A_1 \oplus D_4 \supset A_1 \oplus A_1 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1: \text{non-surviving.} \]

This chain may be discarded due to the sextet/triplet generating rules since it exhibits only two sextets \( (d_3 = 12). \)

1.7.3.1 \( \cdots \supset (A_1)_{12} \oplus A_1 \oplus A_1: \) total pairing.

1.7.3.2 \( \cdots \supset (A_1)_{13} \oplus A_1: \text{non-surviving.} \)

This chain may be discarded due to the sextet/triplet generating rules since it exhibits only one multiplet of dimension 12 and one sextet \( (d_3 = 18). \)

1.7.3.1.1 \( \cdots \supset (A_1)_{12} \oplus A_1: \) total pairing.

1.7.3.1.2 \( \cdots \supset (A_1)_{13}: \) total pairing.

1.7.3.3 \( \cdots \supset (A_1)_{23} \oplus A_1: \) distribution of multiplets identical to that from the chain 1.6.3.3.

1.7.3.3.1 \( \cdots \supset (A_1)_{14} \oplus A_1 \oplus A_1: \text{surviving.} \)

1.7.3.3.2 \( \cdots \supset (A_1)_{15}: \) total pairing.

1.7.3.3.3 \( \cdots \supset (A_1)_{23}: \) distribution of multiplets identical to that from the chain 1.6.3.3.

1.7.3.4 \( \cdots \supset (A_1)_{34} \oplus A_1: \text{total pairing.} \)

• Chain 1.7.4:

\[ B_0 \supset D_0 \supset A_1 \oplus A_1 \oplus D_4 \supset A_1 \oplus A_1 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1: \text{non-surviving.} \]

This chain may be discarded due to the sextet/triplet generating rules since it exhibits only two sextets \( (d_3 = 12). \)

1.7.4.1 \( \cdots \supset (A_1)_{12} \oplus A_1 \oplus A_1: \) total pairing.

1.7.4.2 \( \cdots \supset (A_1)_{13} \oplus A_1: \text{non-surviving.} \)

This chain may be discarded due to the sextet/triplet generating rules since it exhibits only three sextets \( (d_3 = 18). \)

1.7.4.1.1 \( \cdots \supset (A_1)_{12} \oplus A_1: \) total pairing.

1.7.4.2.1 \( \cdots \supset (A_1)_{13}: \) total pairing.

1.7.4.3 \( \cdots \supset (A_1)_{14} \oplus A_1 \oplus A_1: \text{surviving.} \)

1.7.4.3.1 \( \cdots \supset (A_1)_{12} \oplus A_1 \oplus A_1: \) total pairing.

1.7.4.3.2 \( \cdots \supset (A_1)_{13} \oplus A_1 \oplus A_1: \) total pairing.

1.7.4.3.3 \( \cdots \supset (A_1)_{14}: \) distribution of multiplets identical to that from the chain 1.6.4.2.
1.7.4.3.4 \( \cdots \supset (A_1)_{23} \oplus A_1 \oplus A_1 \): distribution of multiplets identical to that from the chain 1.6.4.2.
1.7.4.3.5 \( \cdots \supset (A_1)_{24} \oplus A_1 \oplus A_1 \): distribution of multiplets identical with that obtained from the \( \mathfrak{sp}(6) \)-model after the second step.
1.7.4.3.6 \( \cdots \supset (A_1)_{34} \oplus A_1 \oplus A_1 \): total pairing.
1.7.4.4 \( \cdots \supset (A_1)_{34} \oplus A_1 \oplus A_1 \oplus A_1 \): total pairing.
1.7.4.5 \( \cdots \supset (A_1)_{45} \oplus A_1 \oplus A_1 \oplus A_1 \): surviving.
1.7.4.5.1 \( \cdots \supset (A_1)_{12} \oplus A_1 \oplus A_1 \oplus A_1 \): distribution of multiplets identical to that from the chain 1.6.4.2.
1.7.4.5.2 \( \cdots \supset (A_1)_{14} \oplus A_1 \oplus A_1 \): total pairing.
1.7.4.5.3 \( \cdots \supset (A_1)_{15} \oplus A_1 \oplus A_1 \): total pairing.

- Chain 1.7.5:
\( B_6 \supset D_6 \supset A_1 \oplus A_1 \oplus D_4 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \): non-surviving.
This chain may be discarded due to the sextet/triplet generating rules since it exhibits no triplets, sextets etc. \( (d_3 = 0) \).
1.7.5.1 \( \cdots \supset (A_1)_{12} \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \): total pairing.
1.7.5.2 \( \cdots \supset (A_1)_{13} \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \): distribution of multiplets identical to that from the chain 1.7.4.

- Chain 4.3.2:
\( B_6 \supset C_2 \oplus D_4 \supset C_2 \oplus A_1 \oplus C_2 \supset C_2 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \): non-surviving.
This chain may be discarded due to the sextet/triplet generating rules since it exhibits no triplets, sextets etc. \( (d_3 = 0) \).
4.3.2.1 \( \cdots \supset (A_1)_{12} \oplus A_1 \): non-surviving.
This chain may be discarded due to the sextet/triplet generating rules since it exhibits only one multiplet of dimension 12 and one sextet \( (d_3 = 18) \).
4.3.2.1.1 \( \cdots \supset (A_1)_{12} \): does not reproduce the genetic code.
4.3.2.2 \( \cdots \supset (A_1)_{13} \oplus A_1 \): surviving.
4.3.2.2.1 \( \cdots \supset (A_1)_{12} \): does not reproduce the genetic code.
4.3.2.3 \( \cdots \supset (A_1)_{13} \oplus A_1 \)): total pairing.

- Chain 4.3.4:
\( B_6 \supset C_2 \oplus D_4 \supset C_2 \oplus A_1 \oplus C_2 \supset C_2 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \): non-surviving.
This chain may be discarded due to the sextet/triplet generating rules since it exhibits only one multiplet of dimension 12 \( (d_3 = 12) \).
4.3.4.1 \( \cdots \supset (A_1)_{12} \oplus A_1 \oplus A_1 \): non-surviving.
This chain may be discarded due to the sextet/triplet generating rules since it exhibits only three sextets \( (d_3 = 18) \).
4.3.4.1.1 \( \cdots \supset (A_1)_{12} \oplus A_1 \): total pairing.
4.3.4.1.2 \( \cdots \supset (A_1)_{13} \oplus A_1 \): surviving.
4.3.4.1.2.1 \( \cdots \supset (A_1)_{12} \): does not reproduce the genetic code.
4.3.4.2 \( \cdots \supset (A_1)_{13} \oplus A_1 \oplus A_1 \): surviving.
4.3.4.2.1 \( \cdots \supset (A_1)_{12} \oplus A_1 \): total pairing.
4.3.4.2.2 \( \cdots \supset (A_1)_{13} \oplus A_1 \): distribution of multiplets identical to that from the chain 4.3.4.1.2.
4.3.4.2.3 \( \cdots \supseteq (A_1)_{23} \oplus A_1 \): total pairing.
4.3.4.3 \( \cdots \supseteq (A_1)_{23} \oplus A_1 \oplus A_1 \): total pairing.
4.3.4.4 \( \cdots \supseteq (A_1)_{34} \oplus A_1 \oplus A_1 \): non-surviving.
This chain may be discarded due to the sextet/triplet generating rules in view of the fact that the two multiplets of dimension 12 that occur and that must of course be broken are incapable of generating the three sextets and two triplets in the genetic code: they have highest weights (2-3-0) and (0-3-2) and so each of them can only produce two sextets or four triplets or no sextets and triplets at all, which cannot be combined to yield an odd number of sextets.

4.3.4.4.1 \( \cdots \supseteq (A_1)_{12} \oplus A_1 \): distribution of multiplets identical to that from the chain 4.3.4.1.2.
4.3.4.4.2 \( \cdots \supseteq (A_1)_{13} \oplus A_1 \): total pairing.

- Chain 4.4.1:
  \( B_6 \supseteq C_2 \oplus D_4 \supseteq C_2 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \): non-surviving.
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits no triplets, sextets etc. \((d_3 = 0)\).
4.4.1.1 \( \cdots \supseteq (A_1)_{12} \oplus A_1 \oplus A_1 \oplus A_1 \): non-surviving.
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits only two sextets \((d_3 = 12)\).
4.4.1.1.1 \( \cdots \supseteq (A_1)_{12} \oplus A_1 \oplus A_1 \oplus A_1 \): total pairing.
4.4.1.1.2 \( \cdots \supseteq (A_1)_{13} \oplus A_1 \oplus A_1 \): distribution of multiplets identical to that from the chain 4.3.4.1.
4.4.1.1.3 \( \cdots \supseteq (A_1)_{23} \oplus A_1 \oplus A_1 \): distribution of multiplets identical to that from the chain 4.3.4.2.
4.4.1.1.4 \( \cdots \supseteq (A_1)_{34} \oplus A_1 \oplus A_1 \): total pairing.
4.4.1.2 \( \cdots \supseteq (A_1)_{12} \oplus A_1 \oplus A_1 \oplus A_1 \): total pairing.
4.4.1.3 \( \cdots \supseteq (A_1)_{13} \oplus A_1 \oplus A_1 \oplus A_1 \): distribution of multiplets identical to that from the chain 4.3.4.

- Chain 6.2:
  \( B_6 \supseteq A_1 \oplus A_1 \oplus B_4 \supseteq A_1 \oplus A_1 \oplus A_1 \): non-surviving.
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits no triplets, sextets etc. \((d_3 = 0)\).
6.2.1 \( \cdots \supseteq (A_1)_{12} \oplus A_1 \): total pairing.
6.2.2 \( \cdots \supseteq (A_1)_{13} \oplus A_1 \): non-surviving.
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits only one multiplet of dimension 12 and one sextet \((d_3 = 18)\).
6.2.2.1 \( \cdots \supseteq (A_1)_{12} \): does not reproduce the genetic code.

- Chain 6.4:
  \( B_6 \supseteq A_1 \oplus A_1 \oplus B_4 \supseteq A_1 \oplus A_1 \oplus A_1 \oplus A_1 \): non-surviving.
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits no triplets, sextets etc. \((d_3 = 0)\).
6.4.1 \( \cdots \supseteq (A_1)_{12} \oplus A_1 \oplus A_1 \): total pairing.
6.4.2 \( \cdots \supseteq (A_1)_{13} \oplus A_1 \oplus A_1 \): non-surviving.
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits only one multiplet of dimension 12 and one sextet \((d_3 = 18)\).
6.4.2.1 \cdots \supset (A_1)_{12} \oplus A_1$: total pairing.
6.4.2.2 \cdots \supset (A_1)_{13} \oplus A_1$: total pairing.
6.4.2.3 \cdots \supset (A_1)_{23} \oplus A_1$: surviving.

6.4.2.3.1 \cdots \supset (A_1)_{12}: does not reproduce the genetic code.

Summarizing, we have found a total of 9 chains for $B_6$ that survive up to the end of the second phase and must be subjected to the third phase (breaking of the $A_1$-subalgebras).

$D_7 = \mathfrak{so}(14)$

- Chain 2.3.1.1:
  $D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2 \supset A_2 \oplus A_1 \oplus C_2 \supset A_1^{(1)} \oplus A_1 \oplus A_1$: non-surviving.
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits only one sextet and two triplets ($d_3 = 12$).

- Chain 2.3.1.2:
  $D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2 \supset A_2 \oplus A_1 \oplus C_2 \supset A_1^{(2)} \oplus A_1 \oplus A_1$: surviving.

- Chain 2.3.1.3:
  $D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2 \supset A_2 \oplus A_1 \oplus C_2 \supset A_1^{(1)} \oplus A_1 \oplus A_1$: non-surviving.
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits only one sextet and two triplets ($d_3 = 12$).
2.3.1.3.3 \( \supset (A_1)_{23} \oplus A_1 \oplus A_1 \): total pairing, as well as 4 triplets and 4 singlets.
2.3.1.3.4 \( \supset (A_1)_{34} \oplus A_1 \oplus A_1 \): 4 triplets and 4 singlets.

- Chain 2.3.1.4:
  \( D_7 \supset A_1 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2 \supset A_2 \oplus A_1 \oplus C_2 \supset A_1^{(2)} \oplus A_1 \oplus A_1 \oplus A_1 \): surviving.
  2.3.1.4.1 \( \supset (A_1)_{12} \oplus A_1 \oplus A_1 \): 1 quintet, 3 triplets and 2 singlets.
  2.3.1.4.2 \( \supset (A_1)_{13} \oplus A_1 \oplus A_1 \): surviving.
    2.3.1.4.2.1 \( \supset (A_1)_{12} \oplus A_1 \): total pairing, as well as 2 quintets, 6 triplets and 4 singlets.
    2.3.1.4.2.2 \( \supset (A_1)_{13} \oplus A_1 \): 1 nonet, 1 quintet, 5 triplets and 3 singlets.
    2.3.1.4.2.3 \( \supset (A_1)_{23} \oplus A_1 \): total pairing, as well as 2 nonets, 4 triplets and 2 singlets.
  2.3.1.4.3 \( \supset (A_1)_{23} \oplus A_1 \oplus A_1 \): total pairing, as well as 2 nonets, 4 triplets and 2 singlets.
  2.3.1.4.4 \( \supset (A_1)_{34} \oplus A_1 \oplus A_1 \): 2 nonets, 4 triplets and 2 singlets.

- Chain 2.3.3.1:
  \( D_7 \supset A_1 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \): non-surviving. This chain may be discarded due to the sextet/triplet generating rules since it exhibits no triplets, sextets etc. \((d_3 = 0)\).
  2.3.3.1.1 \( \supset (A_1)_{12} \oplus A_1 \oplus A_1 \): distribution of multiplets identical to that from the chain 2.3.3.1.2.
  2.3.3.1.2 \( \supset (A_1)_{13} \oplus A_1 \oplus A_1 \): non-surviving. This chain may be discarded due to the sextet/triplet generating rules in view of the fact that the multiplet of dimension 24 that occurs and that must of course be broken is incapable of generating the three sextets and two triplets in the genetic code: it has highest weight \((2\text{-}1\text{-}3)\) and so can only produce four sextets or eight triplets or no sextets and triplets at all.
    2.3.3.1.2.1 \( \supset (A_1)_{12} \oplus A_1 \): 1 multiplet of dimension 15, 2 quintets, 2 triplets and 1 singlet.
    2.3.3.1.2.2 \( \supset (A_1)_{13} \oplus A_1 \): total pairing.
    2.3.3.1.2.3 \( \supset (A_1)_{23} \oplus A_1 \): distribution of multiplets identical with that obtained from the \( g_2 \) model after the first step.
      2.3.3.1.2.3.1 \( \supset (A_1)_{12} \): does not reproduce the genetic code.
  2.3.3.1.3 \( \supset (A_1)_{14} \oplus A_1 \oplus A_1 \): non-surviving. This chain may be discarded due to the sextet/triplet generating rules in view of the fact that the three multiplets of dimension 12 that occur and that must of course be broken are incapable of generating the three sextets and two triplets in the genetic code: they have highest weights \((5\text{-}1\text{-}0), (2\text{-}1\text{-}1) \text{ and } (1\text{-}1\text{-}2)\) and so the first can only produce two sextets or no sextets and triplets at all whereas each of the other two can only produce two sextets or four triplets or no sextets and triplets at all, which cannot be combined to yield an odd number of sextets.
    2.3.3.1.3.1 \( \supset (A_1)_{12} \oplus A_1 \): 1 nonet, 1 septet, 2 quintets and 2 triplets.
    2.3.3.1.3.2 \( \supset (A_1)_{13} \oplus A_1 \): total pairing.
    2.3.3.1.3.3 \( \supset (A_1)_{23} \oplus A_1 \): distribution of multiplets identical to that from the chain 2.3.3.1.2.3.
  2.3.3.1.4 \( \supset (A_1)_{34} \oplus A_1 \oplus A_1 \): total pairing.
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Chain 2.3.3.2:

\[ D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 : \text{non-surviving.} \]

This chain may be discarded due to the sextet/triplet generating rules since it exhibits only one multiplet of dimension 12 \((d_3 = 12)\).

2.3.3.2.1 \( \rightarrow (A_1)_{12} \oplus A_1 \oplus A_1 \oplus A_1 \): distribution of multiplets identical to that from the chain 2.3.1.4.

2.3.3.2.2 \( \rightarrow (A_1)_{13} \oplus A_1 \oplus A_1 \oplus A_1 \): non-surviving.

This chain may be discarded due to the sextet/triplet generating rules in view of the fact that the two multiplets of dimension 12 that occur and that must of course be broken are incapable of generating the three sextets and two triplets in the genetic code: they have highest weights \((2-1-1-0)\) and \((2-1-0-1)\) and so each of them can only produce two sextets or four triplets or no sextets and triplets at all, which cannot be combined to yield an odd number of sextets.

2.3.3.2.2.1 \( \rightarrow (A_1)_{12} \oplus A_1 \oplus A_1 : 1 \text{ quintet, 3 triplets and 2 singlets.} \)

2.3.3.2.2.2 \( \rightarrow (A_1)_{13} \oplus A_1 \oplus A_1 : \text{total pairing.} \)

2.3.3.2.2.3 \( \rightarrow (A_1)_{23} \oplus A_1 \oplus A_1 : \text{surviving.} \)

2.3.3.2.2.4 \( \rightarrow (A_1)_{34} \oplus A_1 \oplus A_1 : \text{non-surviving.} \)

This chain may be discarded due to the sextet/triplet generating rules in view of the fact that the three multiplets of dimension 12 that occur and that must of course be broken are incapable of generating the three sextets and two triplets in the genetic code: they have highest weights \((2-1-1-0)\), \((2-1-0-1)\) (twice) and so each of them can only produce two sextets or four triplets or no sextets and triplets at all, which cannot be combined to yield an odd number of sextets.

2.3.3.2.2.4.1 \( \rightarrow (A_1)_{12} \oplus A_1 : \text{total pairing.} \)

2.3.3.2.2.4.2 \( \rightarrow (A_1)_{13} \oplus A_1 : 2 \text{ nonets, 4 triplets and 2 singlets.} \)

2.3.3.2.2.4.3 \( \rightarrow (A_1)_{23} \oplus A_1 : 1 \text{ nonet, 1 quintet, 5 triplets and 3 singlets.} \)

2.3.3.2.3 \( \rightarrow (A_1)_{14} \oplus A_1 \oplus A_1 \oplus A_1 : \text{non-surviving.} \)

This chain may be discarded due to the sextet/triplet generating rules in view of the fact that the three multiplets of dimension 12 that occur and that must of course be broken are incapable of generating the three sextets and two triplets in the genetic code: they have highest weights \((2-1-1-0)\), \((2-1-0-1)\) and \((1-1-2-0)\) and so each of them can only produce two sextets or four triplets or no sextets and triplets at all, which cannot be combined to yield an odd number of sextets.

2.3.3.2.3.1 \( \rightarrow (A_1)_{12} \oplus A_1 : \text{distribution of multiplets identical to that from the chain 2.3.1.4.2.} \)

2.3.3.2.3.2 \( \rightarrow (A_1)_{13} \oplus A_1 \oplus A_1 : \text{total pairing.} \)

2.3.3.2.3.3 \( \rightarrow (A_1)_{14} \oplus A_1 \oplus A_1 : \text{distribution of multiplets identical to that from the chain 2.3.3.2.2.4.} \)

2.3.3.2.3.4 \( \rightarrow (A_1)_{23} \oplus A_1 \oplus A_1 : \text{distribution of multiplets identical to that from the chain 2.3.3.2.2.3.} \)
Chain 2.4.1.1:

\[ D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \]  
\[ \supset A_1^{(1)} \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 : \text{non-surviving.} \]

This chain may be discarded due to the sextet/triplet generating rules since it exhibits no triplets, sextets etc. \((d_3 = 0)\).

2.4.1.1.1 \( \supset (A_1)_{12} \oplus A_1 \oplus A_1 \oplus A_1 \) : non-surviving.

This chain may be discarded due to the sextet/triplet generating rules since it exhibits only two sextets \((d_3 = 12)\).

2.4.1.1.2 \( \supset (A_1)_{13} \oplus A_1 \oplus A_1 : \text{total pairing} \).

2.4.1.1.3 \( \supset (A_1)_{23} \oplus A_1 \oplus A_1 : \text{distribution of multiplets identical to that from the chain 2.3.1.3.} \)

2.4.1.1.4 \( \supset (A_1)_{34} \oplus A_1 \oplus A_1 : \text{total pairing}. \)

2.4.1.1.5 \( \supset (A_1)_{23} \oplus A_1 \oplus A_1 \oplus A_1 : \text{distribution of multiplets identical to that from the chain 2.3.1.3.} \)

Chain 2.4.1.2:

\[ D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \]  
\[ \supset A_1^{(2)} \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 : \text{non-surviving}. \]

This chain may be discarded due to the sextet/triplet generating rules in view of the fact that the four multiplets of dimension 12 that occur and that must of course be broken are incapable of generating the three sextets and two triplets in the genetic code: they have highest weights \((2-1-1-0)\) and \((0-1-1-2)\) and so each of them can only produce two sextets or four triplets or no sextets and triplets at all, which cannot be combined to yield an odd number of sextets.

2.4.1.2.1 \( \supset (A_1)_{12} \oplus A_1 \oplus A_1 \oplus A_1 : \text{non-surviving} \).

This chain may be discarded due to the sextet/triplet generating rules in view of the fact that the two multiplets of dimension 12 that occur and that must of course be broken are incapable of generating the three sextets and two triplets in the genetic code: they have highest weights \((2-1-1-0)\) and \((0-1-1-2)\) and so each of them can only produce two sextets or four triplets or no sextets and triplets at all, which cannot be combined to yield an odd number of sextets.
fact that the two multiplets of dimension 12 that occur and that must of course be
broken are incapable of generating the three sextets and two triplets in the genetic code:
they have highest weights (2-1-1-0) and (2-1-0-1) and so each of them can only produce
two sextets or four triplets or no sextets and triplets at all, which cannot be combined
to yield an odd number of sextets.

2.4.1.2.1.1 \cdots \supset (A_1)_{12} \oplus A_1 \oplus A_1: \text{total pairing.}
2.4.1.2.1.2 \cdots \supset (A_1)_{13} \oplus A_1 \oplus A_1: \text{1 quintet, 3 triplets and 2 singlets.}
2.4.1.2.1.3 \cdots \supset (A_1)_{23} \oplus A_1 \oplus A_1: \text{distribution of multiplets identical to that from}
\text{the chain 2.3.1.4.2.}
2.4.1.2.1.4 \cdots \supset (A_1)_{34} \oplus A_1 \oplus A_1: \text{total pairing.}

2.4.1.2.2 \cdots \supset (A_1)_{23} \oplus A_1 \oplus A_1 \oplus A_1: \text{total pairing.}
2.4.1.2.3 \cdots \supset (A_1)_{34} \oplus A_1 \oplus A_1 \oplus A_1: \text{distribution of multiplets identical to that from the}
\text{chain 2.3.1.4.}

- Chain 2.4.3:

\[ D_7 \supset A_4 \oplus D_3 \supset A_4 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1: \text{non-surviving.} \]
This chain may be discarded due to the sextet/triplet generating rules since it exhibits
no triplets, sextets etc. \((d_3 = 0)\).

2.4.3.1 \cdots \supset (A_1)_{12} \oplus A_1 \oplus A_1 \oplus A_1: \text{distribution of multiplets identical to that}
\text{from the chain 2.4.1.2.}
2.4.3.2 \cdots \supset (A_1)_{13} \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1: \text{non-surviving.} \]
This chain may be discarded due to the sextet/triplet generating rules in view of the fact
that the two multiplets of dimension 12 that occur and that must of course be broken
are incapable of generating the three sextets and two triplets in the genetic code: they
have highest weights (2-1-0-1-0) and (2-1-0-0-1) and so each of them can only produce
two sextets or four triplets or no sextets and triplets at all, which cannot be combined
to yield an odd number of sextets.

2.4.3.2.1 \cdots \supset (A_1)_{12} \oplus A_1 \oplus A_1 \oplus A_1: \text{distribution of multiplets identical to that from}
\text{the chain 2.4.1.2.1.}
2.4.3.2.2 \cdots \supset (A_1)_{13} \oplus A_1 \oplus A_1 \oplus A_1: \text{total pairing.}
2.4.3.2.3 \cdots \supset (A_1)_{14} \oplus A_1 \oplus A_1 \oplus A_1: \text{distribution of multiplets identical to that from the}
\text{chain 2.3.3.2.2.}
2.4.3.2.4 \cdots \supset (A_1)_{23} \oplus A_1 \oplus A_1 \oplus A_1: \text{non-surviving.} \]
This chain may be discarded due to the sextet/triplet generating rules in view of the fact
that the four multiplets of dimension 12 that occur and that must of course be broken
are incapable of generating the three sextets and two triplets in the genetic code: they
have highest weights (2-1-1-0), (2-1-0-1), (1-2-1-0) and (1-2-0-1) and so each of them can only produce
two sextets or four triplets or no sextets and triplets at all, which cannot be combined
to yield an odd number of sextets.

2.4.3.2.4.1 \cdots \supset (A_1)_{12} \oplus A_1 \oplus A_1: \text{total pairing.}
2.4.3.2.4.2 \cdots \supset (A_1)_{13} \oplus A_1 \oplus A_1: \text{distribution of multiplets identical to that}
\text{from the chain 2.3.3.2.2.3.}
2.4.3.2.5 \cdots \supset (A_1)_{24} \oplus A_1 \oplus A_1 \oplus A_1: \text{surviving.} \]
2.4.3.2.5.1 \cdots \supset (A_1)_{12} \oplus A_1 \oplus A_1: \text{1 quintet, 3 triplets and 2 singlets.}
2.4.3.2.5.2 \cdots \supset (A_1)_{13} \oplus A_1 \oplus A_1: \text{distribution of multiplets identical to that}
\text{from the chain 2.3.3.2.2.3.}
2.4.3.2.5.3 \cdots \supset (A_1)_{14} \oplus A_1 \oplus A_1: \text{total pairing.}
2.4.3.2.5.4 \cdots \supset (A_1)_{34} \oplus A_1 \oplus A_1$: distribution of multiplets identical to that from the chain 2.3.3.2.3.

2.4.3.2.6 \cdots \supset (A_1)_{34} \oplus A_1 \oplus A_1 \oplus A_1$: distribution of multiplets identical to that from the chain 2.3.3.2.

2.4.3.2.7 \cdots \supset (A_1)_{45} \oplus A_1 \oplus A_1 \oplus A_1$: total pairing.

2.4.3.3 \cdots \supset (A_1)_{34} \oplus A_1 \oplus A_1 \oplus A_1$: total pairing.

2.4.3.4 \cdots \supset (A_1)_{35} \oplus A_1 \oplus A_1 \oplus A_1$: distribution of multiplets identical to that from the chain 2.3.3.2.

- Chain 3.2:
  \[ D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1: \text{non-surviving}. \]
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits only one multiplet of dimension 12 \((d_3 = 12)\).

3.2.1 \cdots \supset (A_1)_{12}: \text{does not reproduce the genetic code.}

- Chain 3.5.2.2:
  \[ D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1 \oplus A_1 \oplus B_3 \supset A_1 \oplus A_1 \oplus A_1 \oplus G_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1: \text{non-surviving}. \]
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits no triplets, sextets etc. \((d_3 = 0)\).

3.5.2.2.1 \cdots \supset (A_1)_{12} \oplus A_1 \oplus A_1: \text{surviving}. \]

3.5.2.2.2 \cdots \supset (A_1)_{14} \oplus A_1 \oplus A_1: \text{non-surviving.} \]

This chain may be discarded due to the sextet/triplet generating rules in view of the fact that the two multiplets of dimension 12 that occur and that must of course be broken are incapable of generating the three sextets and two triplets in the genetic code: they have highest weights \((5-1-0)\) and \((5-0-1)\) and so each of them can only produce two sextets or no sextets and triplets at all, that is, no triplets.

3.5.2.2.2.1 \cdots \supset (A_1)_{12} \oplus A_1 \oplus A_1: \text{1 nonet, 2 septets, 1 quintet, 1 triplet and 1 singlet.} \]

3.5.2.2.2.2 \cdots \supset (A_1)_{23} \oplus A_1: \text{total pairing,} \]

3.5.2.2.3 \cdots \supset (A_1)_{12} \oplus A_1 \oplus A_1: \text{total pairing,} \]

3.5.2.2.4 \cdots \supset (A_1)_{12} \oplus A_1 \oplus A_1: \text{non-surviving,} \]

This chain may be discarded due to the sextet/triplet generating rules since it exhibits only one multiplet of dimension 12 \((d_3 = 12)\).

3.5.2.2.4.1 \cdots \supset (A_1)_{12} \oplus A_1: \text{1 nonet, 2 septets, 1 quintet, 1 triplet and 1 singlet.} \]

3.5.2.2.4.2 \cdots \supset (A_1)_{13} \oplus A_1: \text{total pairing,} \]

3.5.2.2.4.3 \cdots \supset (A_1)_{12} \oplus A_1: \text{surviving.} \]

3.5.2.2.4.3.1 \cdots \supset (A_1)_{12}: \text{does not reproduce the genetic code.} \]

- Chain 3.5.2.3:
  \[ D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1 \oplus A_1 \oplus B_3 \supset A_1 \oplus A_1 \oplus A_1 \oplus G_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1: \text{distribution of multiplets identical to that from the chain 2.4.3.2.} \]

- Chain 4.2.1:
  \[ D_7 \supset C_2 \oplus B_4 \supset C_2 \oplus A_1 \oplus A_1 \oplus A_1: \text{non-surviving.} \]
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits no triplets, sextets etc. \((d_3 = 0)\).

4.2.1.1 \cdots \supset (A_1)_{12}: \text{does not reproduce the genetic code.} \]
• Chain 4.4.1:
  \[ D_7 \supset C_2 \oplus B_1 \supset C_2 \oplus A_1 \oplus A_1 \supset A_1 \oplus A_1 \oplus A_1: \text{non-surviving.} \]
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits no triplets, sextets etc. \((d_3 = 0)\).
  4.4.1.1 \( \cdots \supset (A_1)_{12} \oplus A_1: \text{non-surviving}. \)
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits only one multiplet of dimension 12 and one sextet \((d_3 = 18)\).
  4.4.1.1.1 \( \cdots \supset (A_1)_{12}: \text{does not reproduce the genetic code.} \)
  4.4.1.2 \( \cdots \supset (A_1)_{23} \oplus A_1: \text{total pairing.} \)

• Chain 4.5.1:
  \[ D_7 \supset C_2 \oplus B_4 \supset C_2 \oplus A_1 \oplus A_1 \oplus C_2 \supset (C_2)_{14} \oplus A_1 \oplus A_1 \supset A_1 \oplus A_1 \oplus A_1: \text{non-surviving.} \]
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits only two sextets \((d_3 = 12)\).
  4.5.1.1 \( \cdots \supset (A_1)_{12} \oplus A_1: \text{non-surviving.} \)
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits only three sextets \((d_3 = 18)\).
  4.5.1.1.1 \( \cdots \supset (A_1)_{12}: \text{does not reproduce the genetic code.} \)
  4.5.1.2 \( \cdots \supset (A_1)_{23} \oplus A_1: \text{total pairing.} \)

• Chain 4.5.2:
  \[ D_7 \supset C_2 \oplus B_4 \supset C_2 \oplus A_1 \oplus A_1 \oplus C_2 \supset (C_2)_{14} \oplus A_1 \oplus A_1 \supset A_1 \oplus A_1 \oplus A_1: \text{distribution of multiplets identical to that from the } B_6 \text{-chain 1.7.4.5.} \]

• Chain 4.5.3:
  \[ D_7 \supset C_2 \oplus B_4 \supset C_2 \oplus A_1 \oplus A_1 \oplus C_2 \supset C_2 \oplus A_1 \oplus A_1 \oplus A_1 \supset A_1 \oplus A_1 \oplus A_1: \text{non-surviving.} \]
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits no triplets, sextets etc. \((d_3 = 0)\).
  4.5.3.1 \( \cdots \supset (A_1)_{12} \oplus A_1 \oplus A_1: \text{non-surviving.} \)
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits only one multiplet of dimension 12 \((d_3 = 12)\).
  4.5.3.1.1 \( \cdots \supset (A_1)_{12} \oplus A_1: \text{total pairing.} \)
  4.5.3.1.2 \( \cdots \supset (A_1)_{23} \oplus A_1: \text{distribution of multiplets identical to that from the chain 4.5.1.1.1.} \)
  4.5.3.1.3 \( \cdots \supset (A_1)_{23} \oplus A_1: \text{non-surviving.} \)
  This chain may be discarded due to the sextet/triplet generating rules in view of the fact that the two multiplets of dimension 12 that occur and that must of course be broken are incapable of generating the three sextets and two triplets in the genetic code: they have highest weights \((3-2)\) and \((2-3)\) and so each of them can only produce two sextets or four triplets or no sextets and triplets at all, which cannot be combined to yield an odd number of sextets.
  4.5.3.1.3.1 \( \cdots \supset (A_1)_{12}: \text{does not reproduce the genetic code.} \)
  4.5.3.2 \( \cdots \supset (A_1)_{14} \oplus A_1 \oplus A_1: \text{distribution of multiplets identical to that from the chain 4.5.1.} \)
  4.5.3.3 \( \cdots \supset (A_1)_{23} \oplus A_1 \oplus A_1: \text{total pairing.} \)
• Chain 5.2.2.2:
  \( D_7 \supset B_3 + B_3 \supset B_3 + G_2 \supset G_2 + G_2 \supset G_2 + A_1 \supset A_1 \oplus A_1 \): non-surviving.
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits no triplets, sextets etc. \((d_3 = 0)\).

  5.2.2.2.1 \( \cdots \supset (A_1)_{12} \): does not reproduce the genetic code.

• Chain 5.2.2.3:
  \( D_7 \supset B_3 + B_3 \supset B_3 + G_2 \supset G_2 + G_2 \supset G_2 + A_1 \supset A_1 \oplus A_1 \oplus A_1 \):
distribution of multiplets identical to that from the chain 3.5.2.2.1.

• Chain 5.2.2.4:
  \( D_7 \supset B_3 + B_3 \supset B_3 + G_2 \supset G_2 + G_2 \supset G_2 + A_1 \oplus A_1 \oplus A_1 \oplus A_1 \):
distribution of multiplets identical to that from the chain 3.5.2.2.1.

• Chain 5.2.2.5:
  \( D_7 \supset B_3 + B_3 \supset B_3 + G_2 \supset G_2 + G_2 \supset G_2 + A_1 \oplus A_1 \oplus A_1 \oplus A_1 \):
distribution of multiplets identical to that from the chain 2.4.3.2.5.

• Chain 6.1.1.2:
  \( D_{12} \supset A_1 \oplus A_1 \oplus D_5 \supset A_1 \oplus A_1 \oplus A_4 \supset A_1 \oplus A_1 \oplus A_3 \supset A_1 \oplus A_1 \oplus G_2 \supset A_1 \oplus A_1 \oplus A_1 \): non-surviving.
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits no triplets, sextets etc. \((d_3 = 0)\).

  6.1.1.2.1 \( \cdots \supset (A_1)_{12} \oplus A_1 \): total pairing.
  6.1.1.2.2 \( \cdots \supset (A_1)_{13} \oplus A_1 \): non-surviving.
  This chain may be discarded due to the sextet/triplet generating rules since it exhibits only one sextet and two triplets \((d_4 = 12)\).

  6.1.1.2.1 \( \cdots \supset (A_1)_{12} \): does not reproduce the genetic code.

• Chain 6.1.4.2:
  \( D_{12} \supset A_1 \oplus A_1 \oplus D_3 \supset A_1 \oplus A_1 \oplus A_4 \supset A_1 \oplus A_1 \oplus A_3 \supset A_1 \oplus A_1 \oplus A_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \): non-surviving.
  This chain may be discarded due to the sextet/triplet generating rules in view of the fact that the two multiplets of dimension 12 that occur and that must of course be broken, together with the four sextets that occur and that may or may not be broken, are incapable of generating the three sextets and two triplets in the genetic code: the former have highest weights \((1-0-1-2)\) and \((0-1-1-2)\) and so each of them can only produce two sextets or four triplets or no sextets and triplets at all, whereas the latter are organized into two identical sextets with highest weight \((1-0-0-2)\) plus another two identical sextets with highest weight \((0-1-0-2)\); all this cannot be combined to yield an odd number of sextets.

  6.1.4.2.1 \( \cdots \supset (A_1)_{12} \oplus A_1 \oplus A_1 \): total pairing.
  6.1.4.2.2 \( \cdots \supset (A_1)_{13} \oplus A_1 \oplus A_1 \): surviving.

  6.1.4.2.2.1 \( \cdots \supset (A_1)_{12} \oplus A_1 \): total pairing.
  6.1.4.2.2.2 \( \cdots \supset (A_1)_{13} \oplus A_1 \): 1 quintet, 3 triplets and 2 singlets.
  6.1.4.2.2.3 \( \cdots \supset (A_1)_{23} \oplus A_1 \): surviving.

  6.1.4.2.2.3.1 \( \cdots \supset (A_1)_{12} \): does not reproduce the genetic code.
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6.1.4.2.3 \( \cdots \supset (A_1)_{14} \oplus A_1 \oplus A_1 \): non-surviving.
This chain may be discarded due to the sextet/triplet generating rules in view of the fact that the multiplet of dimension 12 that occurs and that must of course be broken, together with the two sextets that occur and that may or may not be broken, are incapable of generating the three sextets and two triplets in the genetic code: the former has highest weight (2-1-1) and so can only produce two sextets or four triplets or no sextets and triplets at all, whereas the latter are organized into two identical sextets with highest weight (2-1-0); all this cannot be combined to yield an odd number of sextets.

6.1.4.2.3.1 \( \cdots \supset (A_1)_{12} \oplus A_1 \): total pairing.
6.1.4.2.3.2 \( \cdots \supset (A_1)_{13} \oplus A_1 \): 1 quintet, 3 triplets and 2 singlets.
6.1.4.2.3.3 \( \cdots \supset (A_1)_{23} \oplus A_1 \): distribution of multiplets identical to that from the chain 6.1.4.2.2.3.

6.1.4.2.4 \( \cdots \supset (A_1)_{34} \oplus A_1 \oplus A_1 \): non-surviving.
This chain may be discarded due to the sextet/triplet generating rules in view of the fact that the four sextets that occur and that may or may not be broken, are incapable of generating the three sextets and two triplets in the genetic code: they are organized into two identical sextets with highest weight (2-1-0) plus another two identical sextets with highest weight (2-0-1), which cannot be combined to yield an odd number of sextets.

6.1.4.2.4.1 \( \cdots \supset (A_1)_{12} \oplus A_1 \): 1 quintet, 3 triplets and 2 singlets.
6.1.4.2.4.2 \( \cdots \supset (A_1)_{23} \oplus A_1 \): total pairing.

- Chain 6.3.1:
  \( D_7 \supset A_1 \oplus A_1 \oplus D_5 \supset A_1 \oplus A_1 \oplus D_5 \supset A_1 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \): distribution of multiplets identical to that from the chain 3.5.2.2.2.

- Chain 6.3.2:
  \( D_7 \supset A_1 \oplus A_1 \oplus D_5 \supset A_1 \oplus A_1 \oplus D_5 \supset A_1 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \): distribution of multiplets identical to that from the chain 2.4.3.2.4.

Summarizing, we have found a total of 12 chains for \( D_7 \) that survive up to the end of the second phase and must be subjected to the third phase (breaking of the \( A_1 \)-subalgebras).

5. Third Phase of Symmetry Breaking

In the last phase of the symmetry breaking process, we subject the surviving chains from the first two phases to breaking of one or several of the \( A_1 \)-subalgebras, which requires a rather long and tedious case-by-case analysis of each of the corresponding tables (in which the abbreviation “HW” stands for “highest weight”).
$B_6 = \mathfrak{so}(13)$

1. Chain 1.6.3.3:

$B_6 \supset D_6 \supset A_1 \oplus C_3 \supset A_1 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus (A_1)_2 \oplus A_1$

<table>
<thead>
<tr>
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<tr>
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</tbody>
</table>

1 subspace 2 subspaces 4 subspaces 8 subspaces 8 subspaces 10 subspaces

In this chain, the first $\mathfrak{su}(2)$ must be broken, in order to eliminate the multiplets of dimension 10. Now observe that

1. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 20 multiplets with $d_3 = 18$,
2. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 35 multiplets with $d_3 = 12$,
3. breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 13 multiplets with $d_3 = 12$,
4. breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 21 multiplets with $d_3 = 12$.

Note that option 4 leads to a scheme which does not allow for freezing and is remote from the genetic code, with 1 sextet, 4 quintets, 5 quartets, 2 triplets, 3 doublets and 6 singlets. In the case of option 2, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension $> 6$ must not be frozen and will therefore provide 4 triplets and 10 doublets, which is certainly too many.
2. Chain 1.6.4.2:

\[ B_6 \supset D_6 \supset A_1 \oplus C_3 \supset A_1 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus (A_1)^{23} \oplus A_1 \oplus A_1 \]

<table>
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<tr>
<th>$B_6$</th>
<th>$D_6$</th>
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<th>$(A_1)^2 \oplus C_2$</th>
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1 subspace 2 subspaces 4 subspaces 8 subspaces 13 subspaces 15 subspaces

In this chain, breaking the third $\mathfrak{su}(2)$ to $\mathfrak{o}(2)$ has no effect and will be disregarded; moreover, at least one of the three $\mathfrak{su}(2)$'s must be broken down to $\mathfrak{o}(2)$ because otherwise the two octets would remain unbroken. Now observe that

1. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 17 multiplets with $d_3 = 18$ (only two of the five sextets are broken into a quartet and a doublet each).
2. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 25 multiplets with $d_3 = 18$.
3. breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 19 multiplets with $d_3 = 12$ (only three of the five sextets are broken into a quartet and a doublet each and the only quartet is broken into two doublets),
4. breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 30 multiplets with $d_3 = 12$,
5. breaking the third $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates precisely 21 multiplets with $d_3 = 30$.

Note that option 5 leads to an interesting scheme which, without any freezing, comes close to the genetic code but is slightly different, with 3 sextets, 5 quartets, 4 triplets, 5 doublets and 4 singlets. In the cases of options 2 and 4, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, it turns out that there is no possibility of generating, for example, the correct number of singlets (2); we can only get 0 or 4 singlets in the first case and 0, 4, 6 or 10 singlets in the second case, apart from other deviations.
3. Chain 1.7.3.3:

\[ B_6 \supset D_6 \supset A_1 \oplus A_1 \oplus D_4 \supset A_1 \oplus A_1 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \supset (A_1)_{14} \oplus A_1 \oplus A_1 \]

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1 subspace 2 subspaces 4 subspaces 6 subspaces 6 subspaces 8 subspaces

In this chain, the first \( \mathfrak{su}(2) \) must be broken, in order to eliminate the multiplets of dimension 10. Moreover, breaking the second \( \mathfrak{su}(2) \) to \( \mathfrak{o}(2) \) has no effect and will be disregarded. Now observe that, in a first step,

1. breaking the first \( \mathfrak{su}(2) \) down to \( \mathfrak{o}(2) \) generates 17 multiplets with \( d_3 = 12 \),
2. breaking the first \( \mathfrak{su}(2) \) down to \( \mathfrak{so}(2) \) generates 30 multiplets with \( d_3 = 12 \),
3. breaking the second \( \mathfrak{su}(2) \) down to \( \mathfrak{so}(2) \) generates 11 multiplets with \( d_3 = 24 \),
4. breaking the third \( \mathfrak{su}(2) \) down to \( \mathfrak{o}(2) \) generates 10 multiplets with \( d_3 = 12 \),
5. breaking the third \( \mathfrak{su}(2) \) down to \( \mathfrak{so}(2) \) generates precisely 15 multiplets with \( d_3 = 12 \).

In the case of option 2, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the two multiplets of dimension 10 must not be frozen and will therefore provide 10 doublets, which is already one too many. Therefore, the only surviving option for continuing the symmetry breaking process is 3. Hence in a second step,

3.1. breaking the first \( \mathfrak{su}(2) \) down to \( \mathfrak{o}(2) \) generates 20 multiplets with \( d_3 = 12 \),
3.2. breaking the first \( \mathfrak{su}(2) \) down to \( \mathfrak{so}(2) \) generates 40 multiplets with \( d_3 = 12 \),
3.3. breaking the third \( \mathfrak{su}(2) \) down to \( \mathfrak{o}(2) \) generates 14 multiplets with \( d_3 = 12 \),
3.4. breaking the third \( \mathfrak{su}(2) \) down to \( \mathfrak{so}(2) \) generates precisely 21 multiplets with \( d_3 = 12 \).

Note that option 3.4 leads to a scheme which does not allow for freezing and is remote from the genetic code, with 1 sextet, 4 quintets, 5 quartets, 2 triplets, 3 doublets and 6 singlets. In the case of option 3.2, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the two multiplets of dimension 5 must not be frozen and will therefore provide 10 singlets, which is certainly too many.
4. Chain 1.7.4.3:

\[ B_6 \supset D_6 \supset A_1 \oplus A_1 \oplus D_4 \supset A_1 \oplus A_1 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \]

\( \supset (A_1)^4 \oplus A_1 \oplus A_1 \oplus A_1 \)

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1 subspace 2 subspaces 4 subspaces 6 subspaces 10 subspaces 12 subspaces

In this chain, breaking the second and the fourth \( \mathfrak{su}(2) \) to \( \mathfrak{o}(2) \) has no effect and will be disregarded; moreover, at least two of the four \( \mathfrak{su}(2) \)'s must be broken down to \( \mathfrak{o}(2) \) because otherwise at least one of the four octets would remain unbroken. Note also the symmetry of the final distribution of multiplets under simultaneous exchange of the first with the third and the second with the fourth \( \mathfrak{su}(2) \). Now observe that, in a first step,

1. breaking the first or third \( \mathfrak{su}(2) \) down to \( \mathfrak{o}(2) \) generates 14 multiplets with \( d_3 = 12 \) (only two of the four sextets are broken into a quartet and a doublet each),
2. breaking the first or third \( \mathfrak{su}(2) \) down to \( \mathfrak{o}(2) \) generates precisely 21 multiplets with \( d_3 = 12 \),
3. breaking the second or fourth \( \mathfrak{su}(2) \) down to \( \mathfrak{o}(2) \) generates 17 multiplets with \( d_3 = 24 \).

Note that option 2 leads to an interesting scheme which, without any freezing, comes close to the genetic code but is slightly different, with 1 octet, 1 sextet, 6 quartets, 2 triplets, 9 doublets and 2 singlets. Therefore, the only surviving option for continuing the symmetry breaking process is 3; for the sake of definiteness, we choose to break the fourth \( \mathfrak{su}(2) \).

Hence in a second step,

3.1. breaking the first \( \mathfrak{su}(2) \) down to \( \mathfrak{o}(2) \) generates 20 multiplets with \( d_3 = 12 \),
3.2. breaking the first \( \mathfrak{su}(2) \) down to \( \mathfrak{o}(2) \) generates 30 multiplets with \( d_3 = 12 \),
3.3. breaking the second \( \mathfrak{su}(2) \) down to \( \mathfrak{o}(2) \) generates 24 multiplets with \( d_3 = 24 \),
3.4. breaking the third \( \mathfrak{su}(2) \) down to \( \mathfrak{o}(2) \) generates 19 multiplets with \( d_3 = 12 \),
3.5. breaking the third \( \mathfrak{su}(2) \) down to \( \mathfrak{o}(2) \) generates 28 multiplets with \( d_3 = 12 \).

In the cases of options 3.2, 3.3 and 3.5, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. A closer analysis reveals that in all three cases, we are able to produce the correct number of sextets (3), triplets (2) and singlets (2), but there is no possibility of generating the correct number of quartets (5).
and of doublets (9): we can only get 2, 4, 6 or 8 quartets and, correspondingly, 15, 11, 7 or 3 doublets.

5. Chain 1.7.4.5:

\[ B_6 \supset D_6 \supset A_1 \oplus A_1 \oplus D_4 \supset A_1 \oplus A_1 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \]

\[ \supset (A_1)^4 \oplus A_1 \oplus A_1 \oplus A_1 \]

\[
\begin{array}{cccccccccccc}
\text{HW} & d & \text{HW} & d & \text{HW} & d & \text{HW} & d & \text{HW} & d & \text{HW} & d \\
000001 & 64 & 000010 & 32 & 0-1-0010 & 16 & 0-1-0-01 & 10 & 0-1-0-1-1 & 8 & 2-0-1-0 & 6 \\
& & & & & & & & (0-0-1-0) & 2 & & \\
& & & & & & & & & & & \\
& & & & & & & & (0-1-2-0-0) & 6 & (0-1-2-0-0) & 6 & (0-0-1-2) & 6 \\
& & & & & & & & & & & \\
& & & & & & & & (1-0-0001) & 16 & (1-0-1-10) & 16 & (1-0-1-1-0) & 8 & (1-1-0-1) & 8 \\
& & & & & & & & & & & & (1-0-1-0-1) & 8 & (1-1-0-1) & 8 \\
& & & & & & & & & & & \\
& & & & & & & & (000001) & 32 & (0-1-0001) & 16 & (0-1-1-10) & 16 & (0-1-1-1-0) & 8 & (1-0-1-1) & 8 \\
& & & & & & & & & & & & (0-1-1-0-1) & 8 & (1-1-0-1) & 8 \\
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& & & & & & & & (1-0-2-0-0) & 6 & (1-0-2-0-0) & 6 & (0-1-0-2) & 6 \\
\end{array}
\]

1 subspace 2 subspaces 4 subspaces 6 subspaces 10 subspaces 12 subspaces

In this chain, breaking the second and the third $\text{su}(2)$ to $\text{o}(2)$ has no effect and will be disregarded. Note also the symmetry of the final distribution of multiplets under exchange of the second with the third and the first with the fourth $\text{su}(2)$. Now observe that

1. breaking the second or third $\text{su}(2)$ down to $\text{o}(2)$ generates 18 multiplets with $d_3 = 24$,
   among which there are already 4 triplets and 4 singlets,
2. breaking the first or fourth $\text{su}(2)$ down to $\text{o}(2)$ generates 14 multiplets with $d_3 = 12$
   (only two of the four sextets are broken into a quartet and a doublet each),
3. breaking the first or fourth $\text{su}(2)$ down to $\text{o}(2)$ generates 20 multiplets with $d_3 = 12$.

Therefore, there is no surviving option for continuing the symmetry breaking process.
6. Chain 4.3.2.2 (with one intermediate step omitted):

\[ B_6 \supset C_2 \oplus D_4 \supset C_2 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \supset (A_1)_{13} \oplus A_1 \]

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\begin{itemize}
  \item 1 subspace
  \item 2 subspaces
  \item 3 subspaces
  \item 3 subspaces
  \item 9 subspaces
\end{itemize}

In this chain, the first \( \mathfrak{su}(2) \) must be broken, in order to eliminate the multiplets of dimension 10 and 14. Now observe that

1. breaking the first \( \mathfrak{su}(2) \) down to \( \mathfrak{o}(2) \) generates 22 multiplets with \( d_3 = 12 \),
2. breaking the first \( \mathfrak{su}(2) \) down to \( \mathfrak{so}(2) \) generates 40 multiplets with \( d_3 = 12 \),
3. breaking the second \( \mathfrak{su}(2) \) down to \( \mathfrak{o}(2) \) generates 10 multiplets with \( d_3 = 12 \),
4. breaking the second \( \mathfrak{su}(2) \) down to \( \mathfrak{so}(2) \) generates 15 multiplets with \( d_3 = 12 \).

In the cases of options 1 and 2, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension > 6 must not be frozen. In the case of option 1, we see that all resulting multiplets still have even dimension. In the case of option 2, we get at least 8 singlets and 4 triplets, which is certainly too many.
7. Chain 4.3.4.1.2 (with one intermediate step omitted):
\[ B_6 \supset C_2 \oplus D_4 \supset C_2 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \supset (A_1)_{12} \oplus A_1 \oplus A_1 \supset (A_1)_{23} \oplus A_1 \]

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In this chain, the second su(2) must be broken, in order to eliminate the multiplets of dimension 10. Now observe that

1. breaking the first su(2) down to \( o(2) \) generates 11 multiplets with \( d_3 = 18 \),
2. breaking the first su(2) down to \( so(2) \) generates 16 multiplets with \( d_3 = 18 \),
3. breaking the second su(2) down to \( o(2) \) generates 22 multiplets with \( d_3 = 12 \),
4. breaking the second su(2) down to \( so(2) \) generates 40 multiplets with \( d_3 = 12 \).

In the cases of options 3 and 4, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension \( > 6 \) must not be frozen. In the case of option 3, we see that all resulting multiplets still have even dimension. In the case of option 4, there is no possibility of generating the correct number of triplets (2); we can only get 0 or 4 triplets.

8. Chain 4.3.4.2 (with one intermediate step omitted):
\[ B_6 \supset C_2 \oplus D_4 \supset C_2 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \supset (A_1)_{13} \oplus A_1 \oplus A_1 \]

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1 subspace 2 subspaces 3 subspaces 5 subspaces 7 subspaces
In this chain, the first $su(2)$ must be broken, in order to eliminate the multiplets of dimension 10. Moreover, breaking the third $su(2)$ to $\sigma(2)$ has no effect and will be disregarded. Now observe that, in a first step,

1. breaking the first $su(2)$ down to $\sigma(2)$ generates 16 multiplets with $d_3 = 12$,
2. breaking the first $su(2)$ down to $so(2)$ generates 28 multiplets with $d_3 = 12$,
3. breaking the second $su(2)$ down to $\sigma(2)$ generates 8 multiplets with $d_3 = 12$,
4. breaking the second $su(2)$ down to $so(2)$ generates 12 multiplets with $d_3 = 12$,
5. breaking the third $su(2)$ down to $so(2)$ generates 10 multiplets with $d_3 = 24$.

In the case of option 2, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplet of dimension 12 and the two multiplets of dimension 10 must not be frozen and will therefore provide 4 triplets and 10 doublets, which is certainly too many. Therefore, the only surviving option for continuing the symmetry breaking process is 5. Hence in a second step,

5.1. breaking the first $su(2)$ down to $\sigma(2)$ generates 23 multiplets with $d_3 = 12$,
5.2. breaking the first $su(2)$ down to $so(2)$ generates 40 multiplets with $d_3 = 12$,
5.3. breaking the second $su(2)$ down to $\sigma(2)$ generates 11 multiplets with $d_3 = 12$,
5.4. breaking the second $su(2)$ down to $so(2)$ generates 16 multiplets with $d_3 = 12$.

In the cases of options 5.1 and 5.2, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension $> 6$ and of dimension 5 must not be frozen. In the case of option 5.1, we are able to produce the correct number of sextets (3), triplets (2) and singlets (2), but there is no possibility of generating the correct number of quartets (5) and of doublets (9): we can only get 6 or 7 quartets and, correspondingly, 7 or 5 doublets. In the case of option 5.2, we get at least 10 singlets, which is certainly too many.

9. Chain 6.4.2.3:

$B_6 \supset A_1 \oplus A_1 \oplus B_4 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \supset (A_1)_{13} \oplus A_1 \oplus A_1 \supset (A_1)_{23} \oplus A_1$

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1 subspace 2 subspaces 4 subspaces 6 subspaces 8 subspaces

In this chain, both the first and the second $su(2)$ must be broken, in order to eliminate all multiplets whose dimension is a multiple of 5. Note also the symmetry of the final distribution of multiplets under exchange of the first with the second $su(2)$. On the other hand, observe that
1. breaking the first or second $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 14 multiplets with $d_3 = 18$.
2. breaking the first or the second $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 24 multiplets with $d_3 = 18$.

Therefore, there is no possibility to continue the symmetry breaking process to perform the necessary breaking of the other $\mathfrak{su}(2)$.

$D_7 = \mathfrak{so}(14)$

1. Chain 2.3.1.1.2:

$D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2 \supset A_2 \oplus A_1 \oplus C_2 \supset A_1^{(1)} \oplus A_1 \oplus (A_1)_{13} \oplus A_1$

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In this chain, the first $\mathfrak{su}(2)$ must be broken, in order to eliminate the multiplets of dimension 5 and 10. Now observe that

1. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 23 multiplets with $d_3 = 12$.
2. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 40 multiplets with $d_3 = 12$.
3. breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 15 multiplets with $d_3 = 12$.
4. breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates precisely 21 multiplets with $d_3 = 12$.

Note that option 4 leads to a scheme which does not allow for freezing and is remote from the genetic code, with 1 sextet, 4 quintets, 5 quartets, 2 triplets, 3 doublets and 6 singlets. In the cases of options 1 and 2, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension $> 6$ and of dimension 5 must not be frozen. In the case of option 1, we are able to produce the correct number of sextets (3), triplets (2) and singlets (2), but there is no possibility of generating the correct number of quartets (5) and doublets (9): we get at least 6 quartets and at most 7 doublets. In the case of option 2, the multiplets of dimension 5 will provide at least 10 singlets, which is certainly too many.
2. Chain 2.3.1.2:

$$D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2 \supset A_2 \oplus A_1 \oplus C_2 \supset A_1^{(2)} \oplus A_1 \oplus A_1$$

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| 1 subspace | 2 subspaces | 3 subspaces | 6 subspaces | 6 subspaces |

In this chain, the third $\mathfrak{su}(2)$ must be broken, in order to eliminate the multiplets of dimension 5 and 15. Now observe that, in a first step,

1. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 9 multiplets with $d_3 = 12$,
2. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 12 multiplets with $d_3 = 12$,
3. breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 8 multiplets with $d_3 = 48$,
4. breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 12 multiplets with $d_3 = 48$, among which there are already 8 odd-dimensional multiplets, including 3 triplets and 3 singlets,
5. breaking the third $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 12 multiplets with $d_3 = 51$,
6. breaking the third $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 20 multiplets with $d_3 = 51$, among which there are already 12 odd-dimensional multiplets, including 6 triplets and 5 singlets.

Therefore, the only surviving options for continuing the symmetry breaking process are 3 and 5. Hence in a second step,

3.1. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 12 multiplets with $d_3 = 0$,
3.2. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 16 multiplets with $d_3 = 0$,
3.3. breaking the second $\mathfrak{o}(2)$ down to $\mathfrak{so}(2)$ generates 10 multiplets with $d_3 = 48$, among which there are already 6 odd-dimensional multiplets, including 3 triplets,
3.4. breaking the third $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 14 multiplets with $d_3 = 48$,
3.5. breaking the third $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 22 multiplets with $d_3 = 48$,
5.1. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 18 multiplets with $d_3 = 12$,
5.2. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 24 multiplets with $d_3 = 12$,
5.3. breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 14 multiplets with $d_3 = 48$:
   this gives the same distribution of multiplets as option 3.4,
5.4. breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 20 multiplets with $d_3 = 48$, among which there are already 6 odd-dimensional multiplets, namely 3 triplets and 3 singlets,
5.5. breaking the third $\mathfrak{o}(2)$ down to $\mathfrak{so}(2)$ generates 20 multiplets with $d_3 = 51$, among which there are already 12 odd-dimensional multiplets, including 6 triplets and 5 singlets.

In the cases of options 3.5 and 5.2, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension $> 6$ and of dimension 5 must not be frozen. In the case of option 3.5, the multiplets of
dimension 15 and of dimension 5 will break into 5 triplets and 5 singlets, respectively, so we get at least 6 triplets and 6 singlets. In the case of option 5.2, the multiplet of dimension 9 will break into 3 triplets, so we get at least 3 triplets and 6 odd-dimensional multiplets. We are thus left with a single surviving option for continuing the symmetry breaking process, namely 3.4 = 5.3, which consists in breaking the second and the third \( su(2) \) down to \( o(2) \), generating 14 multiplets with \( d_3 = 48 \). Hence in a third step,

3.4.1. breaking the first \( su(2) \) down to \( o(2) \) generates precisely 21 multiplets with \( d_3 = 0 \),
3.4.2. breaking the first \( su(2) \) down to \( so(2) \) generates 28 multiplets with \( d_3 = 0 \),
3.4.3. breaking the second \( o(2) \) down to \( so(2) \) generates 12 multiplets with \( d_3 = 48 \), among which there are already 8 odd-dimensional multiplets, including 3 triplets and 3 singlets,
3.4.4. breaking the third \( o(2) \) down to \( so(2) \) generates 22 multiplets with \( d_3 = 48 \).

Note that option 3.4.1 leads to a scheme which does not allow for freezing and is remote from the genetic code, with 2 octets, 7 quartets, 8 doublets and 4 singlets. In the cases of options 3.4.2 and 3.4.4, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension \( > 6 \) must not be frozen. In the case of option 3.4.2, we are able to produce the correct number of sextets (3), triplets (2) and singlets (2), but there is no possibility of generating the correct number of quartets (5) and doublets (9); we can only get 8 quartets and, correspondingly, 3 doublets. In the case of option 3.4.4, the multiplets of dimension 12 will break into 2 sextets, so we get at least 5 sextets.

3. Chain 2.3.1.3.2:

\[ D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2 \supset A_2 \oplus A_1 \oplus C_2 \supset A_1^{(1)} \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \]

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1 subspace | 2 subspaces | 3 subspaces | 6 subspaces | 15 subspaces | 17 subspaces
In this chain, breaking the third \(\mathfrak{su}(2)\) to \(\mathfrak{o}(2)\) has no effect and will be disregarded; moreover, at least one of the three \(\mathfrak{su}(2)\)'s must be broken down to \(\mathfrak{o}(2)\) because otherwise the octet would remain unbroken. Now observe that

1. breaking the first \(\mathfrak{su}(2)\) down to \(\mathfrak{o}(2)\) generates 19 multiplets with \(d_3 = 12\) (only two of the three sextets are broken into a quartet and a doublet each),
2. breaking the first \(\mathfrak{su}(2)\) down to \(\mathfrak{o}(2)\) generates 28 multiplets with \(d_3 = 12\).
3. breaking the second \(\mathfrak{su}(2)\) down to \(\mathfrak{o}(2)\) generates 20 multiplets with \(d_3 = 12\) (only one of the three sextets is broken into a quartet and a doublet and the two triplets are broken into a doublet and a singlet each).
4. breaking the second \(\mathfrak{su}(2)\) down to \(\mathfrak{o}(2)\) generates 30 multiplets with \(d_3 = 12\),
5. breaking the third \(\mathfrak{su}(2)\) down to \(\mathfrak{o}(2)\) generates 24 multiplets with \(d_3 = 12\).

In the cases of options 2, 4 and 5, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. A closer analysis reveals that in all three cases, we are able to produce the correct number of sextets (3), triplets (2) and singlets (2), but there is no possibility of generating the correct number of quartets (5) and doublets (9): we can only get 4, 6 or 8 quartets and, correspondingly, 11, 7 or 3 doublets.

4. Chain 2.3.1.4:
\[
D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2 \supset A_2 \oplus A_1 \oplus C_2 \supset A_1^{(2)} \oplus A_1 \oplus A_1
\]

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In this chain, breaking the third or fourth \(\mathfrak{su}(2)\) to \(\mathfrak{o}(2)\) has no effect and will be disregarded; moreover, at least one of the first two \(\mathfrak{su}(2)\)'s must be broken because otherwise the nonet would remain unbroken. Note also the symmetry of the final distribution of multiplets under exchange of the third with the fourth \(\mathfrak{su}(2)\). Now observe that, in a first step,

1. breaking the first \(\mathfrak{su}(2)\) down to \(\mathfrak{o}(2)\) generates 15 multiplets with \(d_3 = 12\),
2. breaking the first \(\mathfrak{su}(2)\) down to \(\mathfrak{o}(2)\) generates 20 multiplets with \(d_3 = 12\),
3. breaking the second \(\mathfrak{su}(2)\) down to \(\mathfrak{o}(2)\) generates 12 multiplets with \(d_3 = 48\),
4. breaking the second \(\mathfrak{su}(2)\) down to \(\mathfrak{o}(2)\) generates 18 multiplets with \(d_3 = 48\), among which there are already 4 triplets and 4 singlets,
5. breaking the third \(\mathfrak{su}(2)\) down to \(\mathfrak{o}(2)\) generates 14 multiplets with \(d_3 = 51\).
Therefore, the only surviving options for continuing the symmetry breaking process are 3 and 5. Hence in a second step,

3.1. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 18 multiplets with $d_3 = 0$,
3.2. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 24 multiplets with $d_3 = 0$,
3.3. breaking the second $\mathfrak{o}(2)$ down to $\mathfrak{so}(2)$ generates 18 multiplets with $d_3 = 48$:
   this gives the same distribution of multiplets as option 4 above,
3.4. breaking the third $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 16 multiplets with $d_3 = 48$,
3.5. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates precisely 21 multiplets with $d_3 = 12$,
3.6. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 28 multiplets with $d_3 = 12$,
3.7. breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 16 multiplets with $d_3 = 48$:
   this gives the same distribution of multiplets as option 3.4 above.
5.4. breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 24 multiplets with $d_3 = 48$,
5.5. breaking the fourth $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 20 multiplets with $d_3 = 51$, among
   which there are already 6 triplets and 5 singlets.

Note that option 5.1 leads to an interesting scheme which, without any freezing, comes close to the genetic code but is slightly different, with 1 octet, 1 sextet, 6 quartets, 2 triplets, 9 doublets and 2 singlets. In the cases of options 3.2, 5.2 and 5.4, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension $> 6$ must not be frozen. In the case of option 3.2, the two multiplets of dimension 12 will break into 3 quartets each, so we get at least 9 quartets and at most 2 sextets. In the cases of options 5.2 and 5.4, the multiplet of dimension 9 will break into 3 triplets, so we get at least 4 triplets and 6 odd-dimensional multiplets. We are thus left with a single surviving option for continuing the symmetry breaking process, namely $3.4 = 5.3$, which consists in breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ and the third $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$, generating 16 multiplets with $d_3 = 48$. Hence in

a third step,

3.4.1. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 24 multiplets with $d_3 = 0$,
3.4.2. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 32 multiplets with $d_3 = 0$,
3.4.3. breaking the second $\mathfrak{o}(2)$ down to $\mathfrak{so}(2)$ generates 24 multiplets with $d_3 = 48$,
3.4.4. breaking the fourth $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 22 multiplets with $d_3 = 48$.

In all cases, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplet of dimension 12 must not be frozen. In the case of option 3.4.1, the multiplet of dimension 12 will break into an octet and a quartet. In the cases of options 3.4.3 and 3.4.4, there remain too many multiplets whose dimension is a multiple of 3 (sextets and triplets), since we still have $d_3 = 48$. Finally, in the case of option 3.4.2, we are able to produce the correct number of sextets (3), triplets (2) and singlets (2), but there is no possibility of generating the correct number of quartets (5) and doublets (9); we can only get 4 quartets and, correspondingly, 11 doublets.
5. Chain 2.3.1.4.2:

\[ D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2 \supset A_2 \oplus A_1 \oplus C_2 \supset A_1^{(2)} \oplus A_1 \oplus A_1 \supset (A_1)_{13} \oplus A_1 \oplus A_1 \]

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1 subspace 2 subspaces 3 subspaces 6 subspaces 10 subspaces 12 subspaces

In this chain, breaking the third \( su(2) \) to \( o(2) \) has no effect and will be disregarded; moreover, at least one of the first two \( su(2) \)’s must be broken because otherwise the nonet would remain unbroken. Now observe that, in a first step,

1. breaking the first \( su(2) \) down to \( o(2) \) generates 17 multiplets with \( d_3 = 12 \),
2. breaking the first \( su(2) \) down to \( so(2) \) generates 28 multiplets with \( d_3 = 12 \),
3. breaking the second \( su(2) \) down to \( o(2) \) generates 14 multiplets with \( d_3 = 24 \) (only the nonet is broken into a sextet and a triplet and one of the two triplets is broken into a doublet and a singlet),
4. breaking the second \( su(2) \) down to \( so(2) \) generates precisely 21 multiplets with \( d_3 = 24 \),
5. breaking the third \( su(2) \) down to \( so(2) \) generates 17 multiplets with \( d_3 = 27 \).

Note that option 4 leads to an interesting scheme which, without any freezing, comes close to the genetic code but is slightly different, with 1 octet, 2 sextets, 4 quartets, 4 triplets, 6 doublets and 4 singlets. In the case of option 2, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplet of dimension 9 must not be frozen and will therefore provide 3 triplets, so we get at least 6 triplets and 6 odd-dimensional multiplets and at most 2 sextets. Therefore, the only surviving options for continuing the symmetry breaking process are 3 and 5. Hence in a second step:

3.1. breaking the first \( su(2) \) down to \( o(2) \) generates 20 multiplets with \( d_3 = 0 \),
3.2. breaking the first \( su(2) \) down to \( so(2) \) generates 32 multiplets with \( d_3 = 0 \),
3.3. breaking the second \( o(2) \) down to \( so(2) \) generates precisely 21 multiplets with \( d_3 = 24 \): this gives the same distribution of multiplets as option 4 above,
3.4. breaking the third \( su(2) \) down to \( so(2) \) generates 19 multiplets with \( d_3 = 27 \),
5.1. breaking the first \( su(2) \) down to \( o(2) \) generates 24 multiplets with \( d_3 = 12 \),
5.2. breaking the first \( su(2) \) down to \( so(2) \) generates 40 multiplets with \( d_3 = 12 \),
5.3. breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 19 multiplets with $d_3 = 27$: this gives the same distribution of multiplets as option 3.4 above.

5.4. breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 28 multiplets with $d_3 = 24$.

In the cases of options 3.2, 5.1, 5.2 and 5.4, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension $> 6$ must not be frozen. In the case of option 3.2, the multiplet of dimension 12 will break into 3 quartets, so we get at most 1 sextet. In the cases of options 5.2 and 5.4, the multiplet of dimension 9 will break into 3 triplets, so we get at least 4 triplets and 6 odd-dimensional multiplets and at most 2 sextets. In the case of option 5.1, we are able to produce the correct number of sextets (3), triplets (2) and singlets (2), but there is no possibility of generating the correct number of quartets (5) and doublets (9): we can only get 4 or 6 quartets and, correspondingly, 11 or 7 doublets. We are thus left with a single surviving option for continuing the symmetry breaking process, namely 3.4 = 5.3, which consists in breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ and the third $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$, generating 19 multiplets with $d_3 = 27$. From this, the distribution of multiplets found in the genetic code could be obtained by breaking the remaining octet into two quartets and one of the quartets into two doublets, but this is impossible since the available quartets come in identical pairs. In other words, in any possible further breaking, the number of quartets will be even and the number of doublets will be 3 mod 4.

6. Chain 2.3.3.2.2.3 (with one intermediate step omitted):

$D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \supset (A_1)_{13} \oplus A_1 \oplus A_1 \oplus A_1 \supset (A_1)_{23} \oplus A_1 \oplus A_1$

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1 subspace 2 subspaces 3 subspaces 5 subspaces 8 subspaces 11 subspaces

In this chain, breaking the third $\mathfrak{su}(2)$ to $\mathfrak{o}(2)$ has no effect and will be disregarded. Now observe that, in a first step,

1. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 14 multiplets with $d_3 = 24$,
2. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 22 multiplets with $d_3 = 24$,
3. breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 15 multiplets with $d_3 = 24$,
4. breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 24 multiplets with $d_3 = 24$,
5. breaking the third $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 15 multiplets with $d_3 = 39$. 
In the cases of options 2 and 4, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension \( d > 6 \) must not be frozen. In both cases, the multiplet of dimension 9 will break into 3 triplets, so we get at least 3 triplets and 6 odd-dimensional multiplets. Therefore, the only surviving options for continuing the symmetry breaking process are 1, 3 and 5. Hence in a second step,

1. breaking the first \( \mathfrak{so}(2) \) down to \( \mathfrak{so}(2) \) generates 22 multiplets with \( d_3 = 24 \),
2. breaking the second \( \mathfrak{su}(2) \) down to \( \mathfrak{su}(2) \) generates 19 multiplets with \( d_3 = 0 \),
3. breaking the second \( \mathfrak{su}(2) \) down to \( \mathfrak{su}(2) \) generates 30 multiplets with \( d_3 = 0 \),
4. breaking the third \( \mathfrak{su}(2) \) down to \( \mathfrak{so}(2) \) generates 19 multiplets with \( d_3 = 24 \),

3.1. breaking the first \( \mathfrak{su}(2) \) down to \( \mathfrak{so}(2) \) generates 19 multiplets with \( d_3 = 0 \),
3.2. breaking the first \( \mathfrak{su}(2) \) down to \( \mathfrak{so}(2) \) generates 30 multiplets with \( d_3 = 0 \),
3.3. breaking the second \( \mathfrak{so}(2) \) down to \( \mathfrak{so}(2) \) generates 24 multiplets with \( d_3 = 24 \),
3.4. breaking the third \( \mathfrak{su}(2) \) down to \( \mathfrak{so}(2) \) generates 20 multiplets with \( d_3 = 24 \),

5. breaking the first \( \mathfrak{su}(2) \) down to \( \mathfrak{so}(2) \) generates 19 multiplets with \( d_3 = 24 \); this gives the same distribution of multiplets as option 1.4 above,
5.2. breaking the first \( \mathfrak{su}(2) \) down to \( \mathfrak{so}(2) \) generates 30 multiplets with \( d_3 = 24 \),
5.3. breaking the second \( \mathfrak{so}(2) \) down to \( \mathfrak{so}(2) \) generates 20 multiplets with \( d_3 = 24 \); this gives the same distribution of multiplets as option 3.4 above,
5.4. breaking the second \( \mathfrak{su}(2) \) down to \( \mathfrak{so}(2) \) generates 32 multiplets with \( d_3 = 24 \).

In the cases of options 1.1, 1.3, 3.2, 3.3, 5.2 and 5.4, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension \( d > 6 \) must not be frozen. In the case of option 1.1, we are able to produce the correct number of sextets (3), triplets (2) and singlets (2), but there is no possibility of generating the correct number of quartets (5) and doublets (9): we get at least 6 quartets and at most 7 doublets. In the cases of options 1.3 and 3.2, the multiplet of dimension 12 will break into 3 quartets, so we get at least 6 quartets and at most 6 doublets. In the case of option 3.3, it suffices to freeze the multiplets coming from the nonet, one of the quartets and one of the triplets: this will reproduce the genetic code, as shown in Table 6 of the next section. In the cases of options 5.2 and 5.4, the multiplet of dimension 9 will break into 3 triplets, so we get at least 3 triplets and 6 odd-dimensional multiplets. We are thus left with two surviving options for continuing the symmetry breaking process, namely 1.4 = 5.1 and 3.4 = 5.3. From either of these, the distribution of multiplets found in the genetic code could be obtained by breaking one of the quartets into two doublets (and, in the first case, the remaining octet into two quartets), but this is impossible since the available quartets come in identical pairs (and since, in the first case, the remaining octet can only be broken into two quartets or four doublets, but not into one quartet and two doublets). In other words, in any possible further breaking, the number of quartets will be even and the number of doublets will be 3 mod 4.
7. Chain 2.3.3.2.3.5 (with one intermediate step omitted):

$$D_7 \cong A_3 \oplus D_4 \cong A_3 \oplus A_1 \oplus C_2 \cong A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus (A_1)_{14} \oplus A_1 \oplus A_1 \oplus A_1 \cong (A_1)_{24} \oplus A_1 \oplus A_1$$

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1 subspace 2 subspaces 3 subspaces 5 subspaces 7 subspaces 10 subspaces

In this chain, at least one of the first two $su(2)$'s must be broken because otherwise the nonet would remain unbroken. Note also the symmetry of the final distribution of multiplets under exchange of the first with the second $su(2)$. Now observe that, in a first step,

1. breaking the first $su(2)$ down to $o(2)$ generates 13 multiplets with $d_3 = 36$,
2. breaking the first $su(2)$ down to $so(2)$ generates 20 multiplets with $d_3 = 36$, among which there are already 4 triplets and 4 singlets,
3. breaking the third $su(2)$ down to $o(2)$ generates 11 multiplets with $d_3 = 39$ (only one of the three multiplets of dimension 12 is broken into an octet and a quartet),
4. breaking the third $su(2)$ down to $so(2)$ generates 16 multiplets with $d_3 = 39$.

Therefore, the only surviving options for continuing the symmetry breaking process are 1, 3 and 4. Hence in a second step,

1.1. breaking the first $o(2)$ down to $so(2)$ generates 20 multiplets with $d_3 = 24$: this gives the same distribution of multiplets as option 2 above,
1.2. breaking the second $su(2)$ down to $o(2)$ generates 17 multiplets with $d_3 = 12$,
1.3. breaking the second $su(2)$ down to $so(2)$ generates 26 multiplets with $d_3 = 12$,
1.4. breaking the third $su(2)$ down to $o(2)$ generates 14 multiplets with $d_3 = 24$,
1.5. breaking the third $su(2)$ down to $so(2)$ generates 20 multiplets with $d_3 = 24$,
3.1. breaking the first $su(2)$ down to $o(2)$ generates 14 multiplets with $d_3 = 24$: this gives the same distribution of multiplets as option 1.4 above,
3.2. breaking the first $su(2)$ down to $so(2)$ generates 22 multiplets with $d_3 = 24$,
3.3. breaking the third $o(2)$ down to $so(2)$ generates 16 multiplets with $d_3 = 39$: this gives the same distribution of multiplets as option 4 above,
4.1. breaking the first $su(2)$ down to $o(2)$ generates 20 multiplets with $d_3 = 24$: this gives the same distribution of multiplets as option 1.5 above,
4.2. breaking the first $su(2)$ down to $so(2)$ generates 32 multiplets with $d_3 = 24$. 
In the cases of options 1.3, 3.2 and 4.2, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension > 6 must not be frozen. In the case of option 1.3, we are able to produce the correct number of sextets (3), triplets (2) and singlets (2), but there is no possibility of generating the correct number of quartets (5) and doublets (9); we can only get 6 or 7 or 8 or 9 quartets and, correspondingly, 7 or 5 or 3 or 1 doublets. In the cases of options 3.2 and 4.2, the multiplet of dimension 9 will break into 3 triplets, so we get at least 4 triplets and 6 odd-dimensional multiplets. We are thus left with two surviving options for continuing the symmetry breaking process, namely 1.4 and 1.5, which consist in breaking the first $\text{su}(2)$ down to $\text{o}(2)$ and the third $\text{su}(2)$ down to $\text{o}(2)$ and $\text{so}(2)$, respectively, generating 14 and 20 multiplets, respectively, with $d_3 = 48$. Hence in a third step,

1.4.1. breaking the first $\text{o}(2)$ down to $\text{so}(2)$ generates 22 multiplets with $d_3 = 24$: this gives the same distribution of multiplets as option 3.2 above,
1.4.2. breaking the second $\text{su}(2)$ down to $\text{o}(2)$ generates 18 multiplets with $d_3 = 0$,
1.4.3. breaking the second $\text{su}(2)$ down to $\text{so}(2)$ generates 28 multiplets with $d_3 = 0$,
1.4.4. breaking the third $\text{o}(2)$ down to $\text{so}(2)$ generates 20 multiplets with $d_3 = 24$: this gives the same distribution of multiplets as option 1.5 above,

1.5.1. breaking the first $\text{o}(2)$ down to $\text{so}(2)$ generates 32 multiplets with $d_3 = 24$: this gives the same distribution of multiplets as option 4.2 above,
1.5.2. breaking the second $\text{su}(2)$ down to $\text{o}(2)$ generates 25 multiplets with $d_3 = 0$,
1.5.3. breaking the second $\text{su}(2)$ down to $\text{so}(2)$ generates 40 multiplets with $d_3 = 0$.

In the cases of options 1.4.1, 1.4.3, 1.5.1 and 1.5.3, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension > 6 must not be frozen. In the case of option 1.4.1 (which may differ from option 3.2 when freezing is taken into account), we are able to produce the correct number of sextets (3), triplets (2) and singlets (2), but there is no possibility of generating the correct number of quartets (5) and doublets (9); since the two multiplets of dimension 8 will break into 2 quartets each and since there are two other quartets that remain unbroken during the last step, we get at least 6 quartets and at most 7 doublets. In the case of option 1.4.3, the multiplet of dimension 12 will break into 3 quartets, so we get at most 1 sextet. In the cases of options 1.5.1 and 1.5.3, it suffices to break one quartet into two doublets and freeze all other multiplets: this will reproduce the genetic code, as shown in Table 7 and Table 8 of the next section.
8. Chain 2.4.3.2.5:

\[ \mathcal{D}_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \] 

\[ \supset (A_1)^{13} \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus (A_1)^{24} \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \]

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1 subspace 2 subspaces 4 subspaces 4 subspaces 6 subspaces 9 subspaces

In this chain, breaking the third and the fourth $\mathfrak{su}(2)$ to $\mathfrak{o}(2)$ has no effect and will be disregarded; moreover, at least two of the four $\mathfrak{su}(2)$'s must be broken down to $\mathfrak{o}(2)$ because otherwise there would be octets left over from the multiplet of dimension 16. Note also the symmetry of the final distribution of multiplets under simultaneous exchange of the first with the second and the third with the fourth $\mathfrak{su}(2)$. Now observe that, in a first step,

1. breaking the first or second $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 12 multiplets with $d_3 = 24$,
2. breaking the first or second $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 18 multiplets with $d_3 = 24$, among which there are already 4 triplets and 4 singlets,
3. breaking the third or fourth $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 12 multiplets with $d_3 = 39$.

Therefore, the only surviving options for continuing the symmetry breaking process are 1 and 3; for the sake of definiteness, we choose to break the second $\mathfrak{su}(2)$ in the first case and the fourth $\mathfrak{su}(2)$ in the second case. Hence in a second step,

1.1. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 16 multiplets with $d_3 = 0$,
1.2. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 24 multiplets with $d_3 = 0$,
1.3. breaking the second $\mathfrak{o}(2)$ down to $\mathfrak{o}(2)$ generates 18 multiplets with $d_3 = 24$: this gives the same distribution of multiplets as option 2 above,
1.4. breaking the third $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 15 multiplets with $d_3 = 24$,
1.5. breaking the fourth $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 16 multiplets with $d_3 = 24$,

3.1. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 15 multiplets with $d_3 = 24$: up to the aforementioned symmetry operation of simultaneously exchanging the first with the second and the third with the fourth $\mathfrak{su}(2)$, this gives the same distribution of multiplets as option 1.4 above,
3.2. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 24 multiplets with $d_3 = 24$,
3.3. breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 16 multiplets with $d_3 = 24$: this gives the same distribution of multiplets as option 1.5 above,
3.4. breaking the second $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 24 multiplets with $d_3 = 24$,
3.5. breaking the third $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 16 multiplets with $d_3 = 39$. 


In the cases of options 1.2, 3.2 and 3.4, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension > 6 must not be frozen. In the case of option 1.2, the multiplet of dimension 16 will break into 2 octets. In the cases of options 3.2 and 3.4, the multiplet of dimension 9 will break into 3 triplets, so we get at least 4 triplets and 6 odd-dimensional multiplets. We are thus left with three surviving options for continuing the symmetry breaking process: 1.4, 1.5 and 3.5. Hence in a third step,

1.4.1. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 20 multiplets with $d_3 = 0$,
1.4.2. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 30 multiplets with $d_3 = 0$,
1.4.3. breaking the second $\mathfrak{o}(2)$ down to $\mathfrak{so}(2)$ generates 24 multiplets with $d_3 = 24$,
1.4.4. breaking the fourth $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 20 multiplets with $d_3 = 24$,

1.5.1. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 20 multiplets with $d_3 = 0$,
1.5.2. breaking the first $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 32 multiplets with $d_3 = 0$,
1.5.3. breaking the second $\mathfrak{o}(2)$ down to $\mathfrak{so}(2)$ generates 24 multiplets with $d_3 = 24$,
1.5.4. breaking the third $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 20 multiplets with $d_3 = 24$: this gives the same distribution of multiplets as option 1.4.4 above,

3.5.1. breaking the first or second $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ generates 20 multiplets with $d_3 = 24$: this gives the same distribution of multiplets as option 1.4.4 above,
3.5.2. breaking the first or second $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$ generates 32 multiplets with $d_3 = 24$.

In the cases of options 1.4.2, 1.4.3, 1.5.2, 1.5.3 and 3.5.2, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension > 6 must not be frozen. In the cases of options 1.4.2 and 1.4.3, the 3 multiplets of dimension 8 will break into 2 quartets each, so we get at least 6 quartets. In the case of option 1.5.2, the 2 multiplets of dimension 8 will break into 2 quartets each and the multiplet of dimension 12 will break into 3 quartets, so we get at least 7 quartets and at most 1 sextet. In the case of option 1.5.3, it suffices to freeze the multiplets coming from the nonet, one of the two quartets and one of the two triplets: this will reproduce the genetic code, as shown in Table 9 of the next section. In the case of option 3.5.2, the multiplet of dimension 9 will break into 3 triplets, so we get at least 4 triplets and 6 odd-dimensional multiplets. Finally, we are still left with a single surviving option for continuing the symmetry breaking process, namely $1.4.4 = 3.3.4 = 3.5.1$, which consists in breaking the first or second $\mathfrak{su}(2)$ down to $\mathfrak{o}(2)$ and the third and fourth $\mathfrak{su}(2)$ down to $\mathfrak{so}(2)$, generating 20 multiplets with $d_3 = 24$. From this, the distribution of multiplets found in the genetic code could be obtained by breaking exactly one of the quartets into two doublets, but this is impossible since the freezing mechanism only allows us to get 0, 2, 4 or 6 quartets and, correspondingly, 19, 15, 11 or 7 doublets.
9. Chain 3.5.2.2.1:

\[
D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1 \oplus A_1 \oplus B_3 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus G_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus (A_1)_{12} \oplus A_1 \oplus A_1
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In this chain, the third su(2) must be broken, in order to eliminate the multiplets of dimension 7, 21 and 28. Moreover, breaking the second su(2) to o(2) has no effect and will be disregarded. Now observe that, in a first step,

1. breaking the first su(2) down to o(2) generates 8 multiplets with \( d = 0 \),
2. breaking the first su(2) down to so(2) generates 10 multiplets with \( d = 0 \),
3. breaking the second su(2) down to so(2) generates 8 multiplets with \( d = 24 \),
4. breaking the third su(2) down to o(2) generates 15 multiplets with \( d = 24 \),
5. breaking the third su(2) down to so(2) generates 24 multiplets with \( d = 24 \).

In the case of option 5, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension > 6 must not be frozen, so that we get at least 8 triplets and 8 singlets. Therefore, the only surviving options for continuing the symmetry breaking process are 3 and 4. Hence in a second step,

3.1. breaking the first su(2) down to o(2) generates 10 multiplets with \( d = 0 \),
3.2. breaking the first su(2) down to so(2) generates 16 multiplets with \( d = 0 \),
3.3. breaking the third su(2) down to o(2) generates 20 multiplets with \( d = 24 \),
3.4. breaking the third su(2) down to so(2) generates 32 multiplets with \( d = 24 \),
4.1. breaking the first su(2) down to o(2) generates 20 multiplets with \( d = 0 \),
4.2. breaking the first su(2) down to so(2) generates 32 multiplets with \( d = 0 \),
4.3. breaking the second su(2) down to so(2) generates 20 multiplets with \( d = 24 \): this gives the same distribution of multiplets as option 3.3 above,
4.4. breaking the third o(2) down to so(2) generates 24 multiplets with \( d = 24 \).

In the cases of options 3.4, 4.2 and 4.4, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension > 6 must not be frozen. In the case of option 3.4, we again get at least 8 triplets and 8 singlets. In the cases of options 4.2 and 4.4, the 3 multiplets of dimension 8 coming from the multiplet of dimension 28 will break into 2 quartets each, so we get at least 6 quartets. We are thus left with a single surviving option for continuing the symmetry breaking process, namely 3.3 = 4.3, which consists in breaking the second su(2) down to so(2) and the third su(2) down to o(2), generating 20 multiplets with \( d = 24 \). From this, the distribution of multiplets found in the genetic code could be obtained by breaking
exactly one of the quartets into two doublets, but this is impossible since the freezing mechanism only allows us to get 0 or 6 quartets and, correspondingly, 19 or 7 doublets.

10. Chain 3.5.2.2.4.3:

\[ D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus A_1 \oplus A_1 \oplus B_3 \supset A_1 \oplus A_1 \oplus A_1 \oplus G_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \]
\[ \supset (A_1)_{24} \oplus A_1 \oplus A_1 \supset (A_1)_{23} \oplus A_1 \]

<table>
<thead>
<tr>
<th></th>
<th></th>
<th>(A_1)^3 \oplus B_3</th>
<th>(A_1)^3 \oplus G_2</th>
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<td>d</td>
<td>HW</td>
<td>d</td>
<td>HW</td>
<td>d</td>
<td>HW</td>
</tr>
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<td></td>
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</tbody>
</table>

1 subspace 1 subspace 2 subspaces 4 subspaces 4 subspaces 5 subspaces 7 subspaces

In this chain, the second \( \mathfrak{su}(2) \) must be broken, in order to eliminate the multiplets of dimension 7 and 21. Now observe that, in a first step,

1. breaking the first \( \mathfrak{su}(2) \) down to \( \mathfrak{o}(2) \) generates 9 multiplets with \( d_3 = 0 \),
2. breaking the first \( \mathfrak{su}(2) \) down to \( \mathfrak{so}(2) \) generates 14 multiplets with \( d_3 = 0 \),
3. breaking the second \( \mathfrak{su}(2) \) down to \( \mathfrak{o}(2) \) generates 18 multiplets with \( d_3 = 24 \),
4. breaking the second \( \mathfrak{su}(2) \) down to \( \mathfrak{so}(2) \) generates 32 multiplets with \( d_3 = 24 \).

In the case of option 4, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension \( > 6 \) must not be frozen, so we get at least 7 triplets and 7 singlets. Therefore, the only surviving option for continuing the symmetry breaking process is 3. Hence in a second step,

3.1. breaking the first \( \mathfrak{su}(2) \) down to \( \mathfrak{o}(2) \) generates 23 multiplets with \( d_3 = 0 \),
3.2. breaking the first \( \mathfrak{su}(2) \) down to \( \mathfrak{so}(2) \) generates 36 multiplets with \( d_3 = 0 \),
3.3. breaking the second \( \mathfrak{o}(2) \) down to \( \mathfrak{so}(2) \) generates 32 multiplets with \( d_3 = 24 \).

In all cases, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. In the case of option 3.1, we are able to produce the correct number of sextets (3), triplets (2) and singlets (2), but there is no possibility of generating the correct number of quartets (5) and doublets (9): we can only get 8 quartets and, correspondingly, 3 doublets. In the cases of options 3.2 and 3.3, it suffices to freeze the multiplets coming from the multiplets of dimension 21, 16, 7, 4 and 3: this will reproduce the genetic code, as shown in Table 10 and Table 11 of the next section.
11. Chain 6.1.4.2.2:

\[ D_7 \supset A_1 \oplus A_1 \oplus D_5 \supset A_1 \oplus A_1 \oplus A_4 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1^{(2)} \]
\[ \supset (A_1)_{13} \oplus A_1 \oplus A_1 \]

<table>
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<tr>
<th>( D_7 )</th>
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<th>((A_1)^2 \oplus A_4)</th>
<th>((A_1)^3 \oplus A_2)</th>
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<td>(d)</td>
<td>HW</td>
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</tr>
<tr>
<td>((0-1-0001))</td>
<td>32</td>
<td>((1-0-0100))</td>
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<td>((1-0-1-10))</td>
<td>12</td>
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</tbody>
</table>

1 subspace 2 subspaces 6 subspaces 12 subspaces 12 subspaces 14 subspaces

In this chain, breaking the second \(su(2)\) to \(so(2)\) has no effect and will be disregarded. Now observe that, in a first step,

1. breaking the first \(su(2)\) down to \(o(2)\) generates 16 multiplets with \(d_3 = 48\),
2. breaking the first \(su(2)\) down to \(so(2)\) generates 24 multiplets with \(d_3 = 48\),
3. breaking the second \(su(2)\) down to \(so(2)\) generates 20 multiplets with \(d_3 = 51\), among which there are already 4 triplets and 4 singlets,
4. breaking the third \(su(2)\) down to \(o(2)\) generates precisely 21 multiplets with \(d_3 = 12\),
5. breaking the third \(su(2)\) down to \(so(2)\) generates 28 multiplets with \(d_3 = 12\).

Note that option 4 leads to an interesting scheme which, without any freezing, comes close to the genetic code but is slightly different, with 1 octet, 1 sextet, 6 quartets, 2 triplets, 9 doublets and 2 singlets. In the cases of options 2 and 5, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension > 6 must not be frozen. In both cases, the multiplet of dimension 9 will break into 3 triplets, so we get at least 4 triplets and 6 odd-dimensional multiplets. Therefore, the only surviving option for continuing the symmetry breaking process is 1. Hence in a second step,

1.1. breaking the first \(o(2)\) down to \(so(2)\) generates 24 multiplets with \(d_3 = 48\),
1.2. breaking the second \(su(2)\) down to \(so(2)\) generates 22 multiplets with \(d_3 = 48\),
1.3. breaking the third \(su(2)\) down to \(o(2)\) generates 24 multiplets with \(d_3 = 0\),
1.4. breaking the third \(su(2)\) down to \(so(2)\) generates 32 multiplets with \(d_3 = 0\).

In all cases, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplet of dimension 12 must not be frozen. In the cases of options 1.1 and 1.2, we get at least 4 sextets. In the case of option 1.3, the
multiplet of dimension 12 will break into an octet and a quartet. In the case of option 1.4, we get at most 4 quartets.

12. Chain 6.1.4.2.2.3:

\[ D_7 \supset A_1 \oplus A_1 \oplus D_5 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 (^{2}(A_1)^{13} \oplus A_1 \oplus A_1 \supset (A_1)^{23} \oplus A_1 \]

<table>
<thead>
<tr>
<th>( D_7 )</th>
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<th>( (A_1)^2 \oplus A_4 )</th>
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<td>d</td>
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<td>&amp;dquo;</td>
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<td>(1-0-0-00)</td>
<td>2</td>
<td>(1-0-0-0)</td>
<td>2</td>
</tr>
</tbody>
</table>

In this chain, observe that, in a first step,

1. breaking the first \( \text{su}(2) \) down to \( \text{o}(2) \) generates 24 multiplets with \( d_3 = 12 \),
2. breaking the first \( \text{su}(2) \) down to \( \text{so}(2) \) generates 40 multiplets with \( d_3 = 12 \),
3. breaking the second \( \text{su}(2) \) down to \( \text{o}(2) \) generates 19 multiplets with \( d_3 = 24 \),
4. breaking the second \( \text{su}(2) \) down to \( \text{so}(2) \) generates 28 multiplets with \( d_3 = 24 \).

In the cases of options 1, 2 and 4, the symmetry breaking process must terminate, and we must take into account the possibility of freezing. However, the multiplets of dimension \( > 6 \) must not be frozen. In the case of option 1, we are able to produce the correct number of sextets (3), triplets (2) and singlets (2), but there is no possibility of generating the correct number of quartets (5) and doublets (9): we can only get 6 or 4 quartets and, correspondingly, 7 or 11 doublets. In the cases of options 2 and 4, the multiplet of dimension 9 will break into 3 triplets, so we get at least 4 triplets and 6 odd-dimensional multiplets. Therefore, the only surviving option for continuing the symmetry breaking process is 3, which already presents the correct number of sextets (3), triplets (2) and singlets (2). However, since the remaining octet can only be broken into 2 quartets or 4 doublets and since the remaining 4 quartets come in two identical pairs, there is no possibility of generating the correct number of quartets (5) and doublets (9): in any possible further breaking, the number of quartets will be even and the number of doublets will be \( 3 \mod 4 \).
6. Results

Having concluded the long and tedious case by case analysis reported in the previous three sections, we proceed to summarize the results.

- There is no symmetry breaking scheme starting out from the simple Lie algebra $B_6 = \mathfrak{so}(13)$ capable of reproducing the degeneracies observed in the genetic code.
- There are altogether 6 symmetry breaking schemes starting out from the simple Lie algebra $D_7 = \mathfrak{so}(14)$ that do reproduce the degeneracies observed in the genetic code.

In what follows, we shall summarize the symmetry breaking schemes based on simple Lie algebras of low or medium rank (i.e. except $\mathfrak{su}(64)$, $\mathfrak{so}(64)$ and $\mathfrak{sp}(64)$) that reproduce the distribution of multiplets observed in the standard genetic code. We begin with the single scheme based on $\mathfrak{sp}(6)$ and the two schemes based on $\mathfrak{g}_2$, which are included not only for the sake of completeness but also because, in the process of typesetting, the corresponding tables in Ref. 9 have been distorted in such a way that it has become impossible to infer which of the multiplets are subject to freezing in the last step.

Some comments about the notation used in the following tables seem in order. First, for the sake of brevity, we shall continue to characterize simple Lie algebras by their Dynkin labels, even though this may not always be the most intuitive manner to understand the embeddings as maximal subalgebras that are involved. For example, the chains

$$D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2 \supset (A_1 \oplus A_1) \oplus A_1 \oplus (A_1 \oplus A_1)$$

and

$$D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus (A_1 \oplus A_1) \oplus (A_1 \oplus A_1) \supset (A_1 \oplus A_1) \oplus (A_1 \oplus A_1) \oplus (A_1 \oplus A_1)$$

and

$$D_7 \supset A_1 \oplus B_5 \supset A_1 \oplus (A_1 \oplus A_1 \oplus B_3)$$

that appear in the following tables are readily understood using the isomorphisms $A_1 \cong \mathfrak{so}(3)$, $A_1 \oplus A_1 \cong \mathfrak{so}(4)$, $C_2 \cong \mathfrak{so}(5)$ and $A_3 \cong \mathfrak{so}(6)$ to rewrite them as

$$\mathfrak{so}(14) \supset \mathfrak{so}(6) \oplus \mathfrak{so}(8) \supset \mathfrak{so}(6) \oplus (\mathfrak{so}(3) \oplus \mathfrak{so}(5))$$

$$\supset (\mathfrak{so}(3) \oplus \mathfrak{so}(3)) \oplus \mathfrak{so}(3) \oplus \mathfrak{so}(4)$$

(4)

and

$$\mathfrak{so}(14) \supset \mathfrak{so}(6) \oplus \mathfrak{so}(8) \supset \mathfrak{so}(6) \oplus (\mathfrak{so}(4) \oplus \mathfrak{so}(4))$$

$$\supset (\mathfrak{so}(3) \oplus \mathfrak{so}(3)) \oplus \mathfrak{so}(4) \oplus \mathfrak{so}(4)$$

(5)

and

$$\mathfrak{so}(14) \supset \mathfrak{so}(3) \oplus \mathfrak{so}(11) \supset \mathfrak{so}(3) \oplus (\mathfrak{so}(4) \oplus \mathfrak{so}(7))$$

(6)
respectively. We also continue to abbreviate the direct sum of \( p \) copies of \( \mathfrak{su}(2) \) by \((A_1)^p\) and the expression “highest weight” by “HW”: in the case of \( A_1 = \mathfrak{su}(2) \), this is \( 2s \) where \( s \) is the “spin” and in the case of \( \mathfrak{o}(2) \) or \( \mathfrak{so}(2) \), it is \( 2m \) where \( m \) is the “magnetic quantum number”. In the third phase, breaking of the \( k \)th \( \mathfrak{su}(2) \) to \( \mathfrak{o}(2) \) and to \( \mathfrak{so}(2) \) will be indicated by the symbol \( L^2_{k,z} \) and \( L^3_{k,z} \), respectively, representing the operator that, as explained in Ref. 9, implements this breaking. Finally, freezing of multiplets will be indicated by shading.

**The \( \mathfrak{sp}(6) \)-chain**

\[
\mathfrak{sp}(6) \supset \mathfrak{sp}(4) \oplus \mathfrak{su}(2) \supset \mathfrak{su}(2) \oplus \mathfrak{su}(2) \oplus \mathfrak{su}(2) \\
\supset \mathfrak{su}(2) \oplus \mathfrak{o}(2) \oplus \mathfrak{su}(2) \supset \mathfrak{su}(2) \oplus \mathfrak{o}(2) \oplus \mathfrak{so}(2)
\] (7)

Table 4. Branching of the codon representation of \( \mathfrak{sp}(6) \) in the chain (7).

<table>
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<tr>
<th>( C_3 )</th>
<th>( C_2 \oplus A_1 )</th>
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<th>( L^2_{k,z} )</th>
<th>( (L^3_{k,z}, L^3_{k,z}) )</th>
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<td>( d )</td>
<td>HW</td>
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<tr>
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<td>2</td>
</tr>
</tbody>
</table>

1 subspace 6 subspaces 14 subspaces 16 subspaces 27 subspaces
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The first $g_2$-chain

$g_2 \supset \mathfrak{su}(2) \oplus \mathfrak{su}(2) \supset \mathfrak{so}(2) \oplus \mathfrak{so}(2) \oplus \mathfrak{su}(2)$  \hspace{1cm} (8)

Table 5. Branching of the codon representation of $g_2$ in the chain (8).

<table>
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<th>$(A_1)^2$</th>
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<th>$(L_{2,z}, L_{1,z})$</th>
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<td>$d$</td>
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<tr>
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<td></td>
<td></td>
<td>(2, ±2) 6 (+2, ±2) 2</td>
</tr>
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<td>4 (+1, ±3) 2</td>
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<td>(0, ±4)</td>
<td>2 (0, ±4) 2</td>
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<tr>
<td></td>
<td></td>
<td>(0, ±2)</td>
<td>2 (0, ±2) 2</td>
</tr>
<tr>
<td>(1, 1)</td>
<td>4</td>
<td>(1, ±1)</td>
<td>4 (+1, ±1) 2</td>
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<td></td>
<td></td>
<td></td>
<td>(1, ±1) 2</td>
</tr>
<tr>
<td>(0, 2)</td>
<td>3</td>
<td>(0, ±2)</td>
<td>2 (0, ±2) 2</td>
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<tr>
<td></td>
<td></td>
<td>(0, 0)</td>
<td>1 (0, 0) 1</td>
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</tbody>
</table>

1 subspace 8 subspaces 17 subspaces 36 subspaces
The second $g_2$-chain

$$g_2 \supset \mathfrak{su}(2) \oplus \mathfrak{su}(2) \supset \mathfrak{su}(2) \oplus \mathfrak{o}(2) \supset \mathfrak{su}(2) \oplus \mathfrak{o}(2)$$  \hspace{1cm} (9)

Table 6. Branching of the codon representation of $g_2$ in the chain (9).

<table>
<thead>
<tr>
<th>$G_2$</th>
<th>$(A_1)^2$</th>
<th>$L_{2,z}^2$</th>
<th>$L_{2,z}$</th>
</tr>
</thead>
<tbody>
<tr>
<td>HW</td>
<td>$d$</td>
<td>HW</td>
<td>$d$</td>
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<tr>
<td>(11)</td>
<td>64</td>
<td>(2, 4)</td>
<td>15</td>
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<tr>
<td>(1, 5)</td>
<td>12</td>
<td>(1, ±3)</td>
<td>4</td>
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<tr>
<td>(2, 2)</td>
<td>9</td>
<td>(2, ±2)</td>
<td>6</td>
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<tr>
<td>(3, 1)</td>
<td>8</td>
<td>(3, ±1)</td>
<td>8</td>
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<tr>
<td>(1, 3)</td>
<td>8</td>
<td>(1, ±3)</td>
<td>4</td>
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<tr>
<td>(0, 4)</td>
<td>5</td>
<td>(0, ±4)</td>
<td>2</td>
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<tr>
<td>(1, 1)</td>
<td>4</td>
<td>(1, ±1)</td>
<td>4</td>
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<td></td>
<td></td>
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<tr>
<td>(0, 2)</td>
<td>3</td>
<td>(0, ±2)</td>
<td>2</td>
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</tbody>
</table>

1 subspace 8 subspaces 17 subspaces 30 subspaces

The $\mathfrak{so}(14)$-chain

$$D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1$$
$$\supset (A_1)^3 \oplus A_1 \oplus A_1 \oplus A_1 \oplus (A_1)^3 \oplus A_1 \oplus A_1 \supset \cdots$$ \hspace{1cm} (10)

Tables 7 and 8 exhibit the branching of the codon representation of $\mathfrak{so}(14)$ along this chain (No. 6 of the previous section): Table 7 presents the first phase (first four
Table 7. Branching of the codon representation of $\mathfrak{s}(14)$ in the chain (10): first two phases.

<table>
<thead>
<tr>
<th>$D_7$</th>
<th>$A_3 \oplus D_4$</th>
<th>$A_3 \oplus A_1 \oplus C_2$</th>
<th>$(A_1)^5$</th>
<th>$(A_1)^4$</th>
<th>$(A_1)^3$</th>
</tr>
</thead>
<tbody>
<tr>
<td>HW</td>
<td>$d$</td>
<td>HW</td>
<td>$d$</td>
<td>HW</td>
<td>$d$</td>
</tr>
<tr>
<td>(0000010)</td>
<td>64 (001,010)</td>
<td>32 (001,001)</td>
<td>20 (1,1,0,1,1)</td>
<td>16 (1,1,1,1)</td>
<td>16 (2,1,1)</td>
</tr>
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<tr>
<td></td>
<td></td>
<td></td>
<td>(1,1,0,0,0)</td>
<td>4 (1,1,0,0)</td>
<td>4 (1,1,0)</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>(001,2,00)</td>
<td>12 (1,1,2,0,0)</td>
<td>12 (3,1,0,0)</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>(100,0001)</td>
<td>32 (100,1,10)</td>
<td>32 (1,1,1,1,1)</td>
<td>16 (2,1,1,0)</td>
<td>12 (2,2,0)</td>
<td>9</td>
</tr>
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</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>(0,1,1,0)</td>
<td>4 (2,0,0)</td>
<td>3</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>(0,0,0)</td>
<td>1</td>
<td></td>
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<td></td>
<td></td>
<td></td>
<td>(1,1,1,0,1)</td>
<td>16 (2,1,0,1)</td>
<td>12 (1,2,1)</td>
</tr>
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<td></td>
<td></td>
<td></td>
<td>(0,1,0,1)</td>
<td>4 (1,0,1)</td>
<td>4</td>
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</tbody>
</table>

Table 8. Branching of the codon representation of $\mathfrak{s}(14)$ in the chain (10): third phase.

<table>
<thead>
<tr>
<th>$(A_1)^3$</th>
<th>$L_{2,2}$</th>
<th>$L_{2,3}$</th>
</tr>
</thead>
<tbody>
<tr>
<td>HW</td>
<td>$d$</td>
<td>HW</td>
</tr>
<tr>
<td>(2,1,1)</td>
<td>12</td>
<td>(2,±1,1)</td>
</tr>
<tr>
<td>(0,1,1)</td>
<td>4</td>
<td>(0,±1,1)</td>
</tr>
<tr>
<td>(1,1,0)</td>
<td>4</td>
<td>(1,±1,0)</td>
</tr>
<tr>
<td>(1,3,0)</td>
<td>8</td>
<td>(1,±3,0)</td>
</tr>
<tr>
<td>(1,1,0)</td>
<td>4</td>
<td>(1,±1,0)</td>
</tr>
<tr>
<td>(2,2,0)</td>
<td>9</td>
<td>(2,±2,0)</td>
</tr>
<tr>
<td>(0,2,0)</td>
<td>3</td>
<td>(0,±2,0)</td>
</tr>
<tr>
<td>(0,0,0)</td>
<td>1</td>
<td>(0,0,0)</td>
</tr>
<tr>
<td>(1,2,1)</td>
<td>12</td>
<td>(1,±2,1)</td>
</tr>
<tr>
<td>(1,0,1)</td>
<td>4</td>
<td>(1,±0,1)</td>
</tr>
</tbody>
</table>

1 subspace 2 subspaces 3 subspaces 5 subspaces 8 subspaces 11 subspaces
columns) and second phase (last three columns) whereas Table 8 shows the third phase. Of course, it is evident from the analysis performed in Sections 3 and 4 that there are many other chains that, after completion of the second phase, lead to the same distribution of multiplets under \((A_1)^3\); they are represented in terms of the flow diagram shown in Fig. 1.

![Flow diagram](image)

Fig. 1. Graphical representation of all so(14) chains leading to the same distribution of multiplets under \((A_1)^3\) as the chain (10) (which is indicated by the double arrows).

**The so(14)-chain**

\[
D_7 \supset A_3 \oplus D_4 \supset A_3 \oplus A_1 \oplus C_2 \supset A_1 \oplus A_1 \oplus A_1 \oplus A_1 \oplus A_1
\supset (A_1)_{14} \oplus A_1 \oplus A_1 \oplus A_1 \oplus (A_1)_{24} \oplus A_1 \oplus A_1 \supset \cdots
\]

(11)

Tables 9–11 exhibit the branching of the codon representation of so(14) along this chain (No. 7 of the previous section): Table 9 presents the first phase (first four columns) and second phase (last three columns) whereas Table 10 shows the first option and Table 11 the second option for the third phase, which differ only in the last step. Of course, it is evident from the analysis performed in Sections 3 and 4 that there are many other chains that, after completion of the second phase, lead to the same distribution of multiplets under \((A_1)^3\); they are represented in terms of the flow diagram shown in Fig. 2.
Table 9. Branching of the codon representation of \( \mathfrak{so}(14) \) in the chain (11): first two phases.

<table>
<thead>
<tr>
<th>( D_7 )</th>
<th>( A_3 \oplus D_4 )</th>
<th>( A_3 \oplus A_1 \oplus C_2 )</th>
<th>( (A_1)^5 )</th>
<th>( (A_1)^4 )</th>
<th>( (A_1)^3 )</th>
</tr>
</thead>
<tbody>
<tr>
<td>HW</td>
<td>( d )</td>
<td>HW</td>
<td>( d )</td>
<td>HW</td>
<td>( d )</td>
</tr>
<tr>
<td>(0000010)</td>
<td>64</td>
<td>(001,0010)</td>
<td>32</td>
<td>(001,001)</td>
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</tr>
<tr>
<td>(001,2,00)</td>
<td>12</td>
<td>(1,1,2,0,0)</td>
<td>12</td>
<td>(1,1,2,0)</td>
<td>12</td>
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</tr>
<tr>
<td>(100,0001)</td>
<td>32</td>
<td>(100,1,10)</td>
<td>32</td>
<td>(1,1,1,1,0)</td>
<td>16</td>
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</tbody>
</table>

1 subspace 2 subspaces 3 subspaces 5 subspaces 7 subspaces 10 subspaces

Fig. 2. Graphical representation of all \( \mathfrak{so}(14) \) chains leading to the same distribution of multiplets under \( (A_1)^3 \) as the chain (11) (which is indicated by the double arrows).