<?xml version="1.0" encoding="UTF-8"?>
<!DOCTYPE ST26SequenceListing PUBLIC "-//WIPO//DTD Sequence Listing 1.3//EN"
"ST26SequenceListing_V1_3.dtd">
<ST26SequenceListing dtdVersion="V1_3" fileName="0030-02011PC_Sequence_Listing.xml"
softwareName="WIPO Sequence" softwareVersion="2.1.2" productionDate="2022-10-04">
<ApplicationIdentification>
<IPOfficeCode>US</IPOfficeCode>
<ApplicationNumberText></ApplicationNumberText>
<FilingDate></FilingDate>
</ApplicationIdentification>
<ApplicantFileReference>0030-02011PC</ApplicantFileReference>
<EarliestPriorityApplicationIdentification>
<IPOfficeCode>US</IPOfficeCode>
<ApplicationNumberText>63/273,411</ApplicationNumberText>
<FilingDate>2021-10-29</FilingDate>
</EarliestPriorityApplicationIdentification>
<ApplicantName languageCode="en">Seagen Inc.</ApplicantName>
<InventionTitle languageCode="en">METHODS OF TREATING CANCER WITH A COMBINATION OF AN
ANTI-PD-1 ANTIBODY AND AN ANTI-CD30 ANTIBODY-DRUG CONJUGATE</InventionTitle>
<SequenceTotalQuantity>34</SequenceTotalQuantity>
<SequenceData sequenceIDNumber="1">
<INSDSeq>
<INSDSeq_length>5</INSDSeq_length>
<INSDSeq_moltype>AA</INSDSeq_moltype>
<INSDSeq_division>PAT</INSDSeq_division>
<INSDSeq_feature-table>
<INSDFeature>
<INSDFeature_key>source</INSDFeature_key>
<INSDFeature_location>1..5</INSDFeature_location>
<INSDFeature_quals>
<INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>
<INSDQualifier id="q1">

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic
construct</INSDQualifier_value>
</INSDQualifier>
</INSDFeature_quals>
</INSDFeature>
</INSDSeq_feature-table>
<INSDSeq_sequence>DYYIT</INSDSeq_sequence>
</INSDSeq>
</SequenceData>
<SequenceData sequenceIDNumber="2">

<INSDSeq>
<INSDSeq_length>17</INSDSeq_length>
<INSDSeq_moltype>AA</INSDSeq_moltype>
<INSDSeq_division>PAT</INSDSeq_division>
<INSDSeq_feature-table>
<INSDFeature>
<INSDFeature_key>source</INSDFeature_key>
<INSDFeature_location>1..17</INSDFeature_location>
<INSDFeature_quals>
<INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>
<INSDQualifier id="q2">

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic
construct</INSDQualifier_value>
</INSDQualifier>
</INSDFeature_quals>
</INSDFeature>
</INSDSeq_feature-table>
<INSDSeq_sequence>WIYPGSGNTKYNEKFKG</INSDSeq_sequence>
</INSDSeq>
</SequenceData>
<SequenceData sequenceIDNumber="3">
<INSDSeq>
<INSDSeq_length>8</INSDSeq_length>
<INSDSeq_moltype>AA</INSDSeq_moltype>
<INSDSeq_division>PAT</INSDSeq_division>
<INSDSeq_feature-table>
<INSDFeature>
<INSDFeature_key>source</INSDFeature_key>
<INSDFeature_location>1..8</INSDFeature_location>
<INSDFeature_quals>
<INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>
<INSDQualifier id="q3">

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic
construct</INSDQualifier_value>
</INSDQualifier>

```
                                        </INSDFeature_quals>
                                    </INSDFeature>
                                </INSDSeq_feature-table>
                                <INSDSeq_sequence>YGNYWFAY</INSDSeq_sequence>
                        </INSDSeq>
                </SequenceData>
                <SequenceData sequenceIDNumber="4">
                        <INSDSeq>
                                <INSDSeq_length>15</INSDSeq_length>
                                <INSDSeq_moltype>AA</INSDSeq_moltype>
                                <INSDSeq_division>PAT</INSDSeq_division>
                                <INSDSeq_feature-table>
                                        <INSDFeature>
                                                <INSDFeature_key>source</INSDFeature_key>
                                                <INSDFeature_location>1..15</INSDFeature_location>
                                                <INSDFeature_quals>
                                                        <INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
                                                        </INSDQualifier>
                                                        <INSDQualifier id="q4">

<INSDQualifier_name>organism</INSDQualifier_name>
                                                                <INSDQualifier_value>synthetic
construct</INSDQualifier_value>
                                                        </INSDQualifier>
                                                </INSDFeature_quals>
                                        </INSDFeature>
                                </INSDSeq_feature-table>
                                <INSDSeq_sequence>KASQSVDFDGDSYMN</INSDSeq_sequence>
                        </INSDSeq>
                </SequenceData>
                <SequenceData sequenceIDNumber="5">
                        <INSDSeq>
                                <INSDSeq_length>7</INSDSeq_length>
                                <INSDSeq_moltype>AA</INSDSeq_moltype>
                                <INSDSeq_division>PAT</INSDSeq_division>
                                <INSDSeq_feature-table>
                                        <INSDFeature>
                                                <INSDFeature_key>source</INSDFeature_key>
                                                <INSDFeature_location>1..7</INSDFeature_location>
                                                <INSDFeature_quals>
                                                        <INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
```

```
                                                    </INSDQualifier>
                                                    <INSDQualifier id="q5">

<INSDQualifier_name>organism</INSDQualifier_name>
                                                            <INSDQualifier_value>synthetic
construct</INSDQualifier_value>
                                                    </INSDQualifier>
                                            </INSDFeature_quals>
                                    </INSDFeature>
                            </INSDSeq_feature-table>
                            <INSDSeq_sequence>AASNLES</INSDSeq_sequence>
                    </INSDSeq>
            </SequenceData>
            <SequenceData sequenceIDNumber="6">
                    <INSDSeq>
                            <INSDSeq_length>9</INSDSeq_length>
                            <INSDSeq_moltype>AA</INSDSeq_moltype>
                            <INSDSeq_division>PAT</INSDSeq_division>
                            <INSDSeq_feature-table>
                                    <INSDFeature>
                                            <INSDFeature_key>source</INSDFeature_key>
                                            <INSDFeature_location>1..9</INSDFeature_location>
                                            <INSDFeature_quals>
                                                    <INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
                                                    </INSDQualifier>
                                                    <INSDQualifier id="q6">

<INSDQualifier_name>organism</INSDQualifier_name>
                                                            <INSDQualifier_value>synthetic
construct</INSDQualifier_value>
                                                    </INSDQualifier>
                                            </INSDFeature_quals>
                                    </INSDFeature>
                            </INSDSeq_feature-table>
                            <INSDSeq_sequence>QQSNEDPWT</INSDSeq_sequence>
                    </INSDSeq>
            </SequenceData>
            <SequenceData sequenceIDNumber="7">
                    <INSDSeq>
                            <INSDSeq_length>117</INSDSeq_length>
                            <INSDSeq_moltype>AA</INSDSeq_moltype>
                            <INSDSeq_division>PAT</INSDSeq_division>
                            <INSDSeq_feature-table>
                                    <INSDFeature>
                                            <INSDFeature_key>source</INSDFeature_key>
```

<INSDFeature_location>1..117</INSDFeature_location>
<INSDFeature_quals>
<INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>
<INSDQualifier id="q7">

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic
construct</INSDQualifier_value>
</INSDQualifier>
</INSDFeature_quals>
</INSDFeature>
</INSDSeq_feature-table>

<INSDSeq_sequence>QIQLQQSGPEVVKPGASVKISCKASGYTFTDYYITWVKQKPGQGLEWIGWIYPGSGNTKYNEKFKGK
ATLTVDTSSSTAFMQLSSLTSEDTAVYFCANYGNYWFAYWGQGTQVTVSA</INSDSeq_sequence>
</INSDSeq>
</SequenceData>
<SequenceData sequenceIDNumber="8">
<INSDSeq>
<INSDSeq_length>111</INSDSeq_length>
<INSDSeq_moltype>AA</INSDSeq_moltype>
<INSDSeq_division>PAT</INSDSeq_division>
<INSDSeq_feature-table>
<INSDFeature>
<INSDFeature_key>source</INSDFeature_key>
<INSDFeature_location>1..111</INSDFeature_location>
<INSDFeature_quals>
<INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>
<INSDQualifier id="q8">

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic
construct</INSDQualifier_value>
</INSDQualifier>
</INSDFeature_quals>
</INSDFeature>
</INSDSeq_feature-table>

<INSDSeq_sequence>DIVLTQSPASLAVSLGQRATISCKASQSVDFDGDSYMNWYQQKPGQPPKVLIYAASNLESGIPARFS

```
GSGSGTDFTLNIHPVEEEDAATYYCQQSNEDPWTFGGGTKLEIK</INSDSeq_sequence>
                </INSDSeq>
        </SequenceData>
        <SequenceData sequenceIDNumber="9">
                <INSDSeq>
                        <INSDSeq_length>30</INSDSeq_length>
                        <INSDSeq_moltype>AA</INSDSeq_moltype>
                        <INSDSeq_division>PAT</INSDSeq_division>
                        <INSDSeq_feature-table>
                                <INSDFeature>
                                        <INSDFeature_key>source</INSDFeature_key>
                                        <INSDFeature_location>1..30</INSDFeature_location>
                                        <INSDFeature_quals>
                                                <INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
                                                </INSDQualifier>
                                                <INSDQualifier id="q9">

<INSDQualifier_name>organism</INSDQualifier_name>
                                                        <INSDQualifier_value>synthetic
construct</INSDQualifier_value>
                                                </INSDQualifier>
                                        </INSDFeature_quals>
                                </INSDFeature>
                        </INSDSeq_feature-table>

<INSDSeq_sequence>QIQLQQSGPEVVKPGASVKISCKASGYTFT</INSDSeq_sequence>
                </INSDSeq>
        </SequenceData>
        <SequenceData sequenceIDNumber="10">
                <INSDSeq>
                        <INSDSeq_length>14</INSDSeq_length>
                        <INSDSeq_moltype>AA</INSDSeq_moltype>
                        <INSDSeq_division>PAT</INSDSeq_division>
                        <INSDSeq_feature-table>
                                <INSDFeature>
                                        <INSDFeature_key>source</INSDFeature_key>
                                        <INSDFeature_location>1..14</INSDFeature_location>
                                        <INSDFeature_quals>
                                                <INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
                                                </INSDQualifier>
                                                <INSDQualifier id="q10">
```

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic construct</INSDQualifier_value>
</INSDQualifier>
</INSDFeature_quals>
</INSDFeature>
</INSDSeq_feature-table>
<INSDSeq_sequence>WVKQKPGQGLEWIG</INSDSeq_sequence>
</INSDSeq>
</SequenceData>
<SequenceData sequenceIDNumber="11">
<INSDSeq>
<INSDSeq_length>32</INSDSeq_length>
<INSDSeq_moltype>AA</INSDSeq_moltype>
<INSDSeq_division>PAT</INSDSeq_division>
<INSDSeq_feature-table>
<INSDFeature>
<INSDFeature_key>source</INSDFeature_key>
<INSDFeature_location>1..32</INSDFeature_location>
<INSDFeature_quals>
<INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>
<INSDQualifier id="q11">

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic construct</INSDQualifier_value>
</INSDQualifier>
</INSDFeature_quals>
</INSDFeature>
</INSDSeq_feature-table>

<INSDSeq_sequence>KATLTVDTSSSTAFMQLSSLTSEDTAVYFCAN</INSDSeq_sequence>
</INSDSeq>
</SequenceData>
<SequenceData sequenceIDNumber="12">
<INSDSeq>
<INSDSeq_length>11</INSDSeq_length>
<INSDSeq_moltype>AA</INSDSeq_moltype>
<INSDSeq_division>PAT</INSDSeq_division>
<INSDSeq_feature-table>
<INSDFeature>
<INSDFeature_key>source</INSDFeature_key>
<INSDFeature_location>1..11</INSDFeature_location>

<INSDFeature_quals>
<INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>
<INSDQualifier id="q12">

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic
construct</INSDQualifier_value>
</INSDQualifier>
</INSDFeature_quals>
</INSDFeature>
</INSDSeq_feature-table>
<INSDSeq_sequence>WGQGTQVTVSA</INSDSeq_sequence>
</INSDSeq>
</SequenceData>
<SequenceData sequenceIDNumber="13">
<INSDSeq>
<INSDSeq_length>23</INSDSeq_length>
<INSDSeq_moltype>AA</INSDSeq_moltype>
<INSDSeq_division>PAT</INSDSeq_division>
<INSDSeq_feature-table>
<INSDFeature>
<INSDFeature_key>source</INSDFeature_key>
<INSDFeature_location>1..23</INSDFeature_location>
<INSDFeature_quals>
<INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>
<INSDQualifier id="q13">

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic
construct</INSDQualifier_value>
</INSDQualifier>
</INSDFeature_quals>
</INSDFeature>
</INSDSeq_feature-table>
<INSDSeq_sequence>DIVLTQSPASLAVSLGQRATISC</INSDSeq_sequence>
</INSDSeq>
</SequenceData>
<SequenceData sequenceIDNumber="14">
<INSDSeq>

<INSDSeq_length>15</INSDSeq_length>
<INSDSeq_moltype>AA</INSDSeq_moltype>
<INSDSeq_division>PAT</INSDSeq_division>
<INSDSeq_feature-table>
<INSDFeature>
<INSDFeature_key>source</INSDFeature_key>
<INSDFeature_location>1..15</INSDFeature_location>
<INSDFeature_quals>
<INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>
<INSDQualifier id="q14">

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic
construct</INSDQualifier_value>
</INSDQualifier>
</INSDFeature_quals>
</INSDFeature>
</INSDSeq_feature-table>
<INSDSeq_sequence>WYQQKPGQPPKVLIY</INSDSeq_sequence>
</INSDSeq>
</SequenceData>
<SequenceData sequenceIDNumber="15">
<INSDSeq>
<INSDSeq_length>32</INSDSeq_length>
<INSDSeq_moltype>AA</INSDSeq_moltype>
<INSDSeq_division>PAT</INSDSeq_division>
<INSDSeq_feature-table>
<INSDFeature>
<INSDFeature_key>source</INSDFeature_key>
<INSDFeature_location>1..32</INSDFeature_location>
<INSDFeature_quals>
<INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>
<INSDQualifier id="q15">

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic
construct</INSDQualifier_value>
</INSDQualifier>
</INSDFeature_quals>

```
                                        </INSDFeature>
                                </INSDSeq_feature-table>

<INSDSeq_sequence>GIPARFSGSGSGTDFTLNIHPVEEEDAATYYC</INSDSeq_sequence>
                        </INSDSeq>
                </SequenceData>
                <SequenceData sequenceIDNumber="16">
                        <INSDSeq>
                                <INSDSeq_length>10</INSDSeq_length>
                                <INSDSeq_moltype>AA</INSDSeq_moltype>
                                <INSDSeq_division>PAT</INSDSeq_division>
                                <INSDSeq_feature-table>
                                        <INSDFeature>
                                                <INSDFeature_key>source</INSDFeature_key>
                                                <INSDFeature_location>1..10</INSDFeature_location>
                                                <INSDFeature_quals>
                                                        <INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
                                                        </INSDQualifier>
                                                        <INSDQualifier id="q16">

<INSDQualifier_name>organism</INSDQualifier_name>
                                                                <INSDQualifier_value>synthetic
construct</INSDQualifier_value>
                                                        </INSDQualifier>
                                                </INSDFeature_quals>
                                        </INSDFeature>
                                </INSDSeq_feature-table>
                                <INSDSeq_sequence>FGGGTKLEIK</INSDSeq_sequence>
                        </INSDSeq>
                </SequenceData>
                <SequenceData sequenceIDNumber="17">
                        <INSDSeq>
                                <INSDSeq_length>5</INSDSeq_length>
                                <INSDSeq_moltype>AA</INSDSeq_moltype>
                                <INSDSeq_division>PAT</INSDSeq_division>
                                <INSDSeq_feature-table>
                                        <INSDFeature>
                                                <INSDFeature_key>source</INSDFeature_key>
                                                <INSDFeature_location>1..5</INSDFeature_location>
                                                <INSDFeature_quals>
                                                        <INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
```

```
                                                    </INSDQualifier>
                                                    <INSDQualifier id="q17">

<INSDQualifier_name>organism</INSDQualifier_name>
                                                            <INSDQualifier_value>synthetic
construct</INSDQualifier_value>
                                                    </INSDQualifier>
                                            </INSDFeature_quals>
                                    </INSDFeature>
                            </INSDSeq_feature-table>
                            <INSDSeq_sequence>NYYMY</INSDSeq_sequence>
                    </INSDSeq>
            </SequenceData>
            <SequenceData sequenceIDNumber="18">
                    <INSDSeq>
                            <INSDSeq_length>17</INSDSeq_length>
                            <INSDSeq_moltype>AA</INSDSeq_moltype>
                            <INSDSeq_division>PAT</INSDSeq_division>
                            <INSDSeq_feature-table>
                                    <INSDFeature>
                                            <INSDFeature_key>source</INSDFeature_key>
                                            <INSDFeature_location>1..17</INSDFeature_location>
                                            <INSDFeature_quals>
                                                    <INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
                                                    </INSDQualifier>
                                                    <INSDQualifier id="q18">

<INSDQualifier_name>organism</INSDQualifier_name>
                                                            <INSDQualifier_value>synthetic
construct</INSDQualifier_value>
                                                    </INSDQualifier>
                                            </INSDFeature_quals>
                                    </INSDFeature>
                            </INSDSeq_feature-table>
                            <INSDSeq_sequence>GINPSNGGTNFNEKFKN</INSDSeq_sequence>
                    </INSDSeq>
            </SequenceData>
            <SequenceData sequenceIDNumber="19">
                    <INSDSeq>
                            <INSDSeq_length>11</INSDSeq_length>
                            <INSDSeq_moltype>AA</INSDSeq_moltype>
                            <INSDSeq_division>PAT</INSDSeq_division>
                            <INSDSeq_feature-table>
                                    <INSDFeature>
                                            <INSDFeature_key>source</INSDFeature_key>
```

<INSDFeature_location>1..11</INSDFeature_location>
<INSDFeature_quals>
<INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>
<INSDQualifier id="q19">

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic construct</INSDQualifier_value>
</INSDQualifier>
</INSDFeature_quals>
</INSDFeature>
</INSDSeq_feature-table>
<INSDSeq_sequence>RDYRFDMGFDY</INSDSeq_sequence>
</INSDSeq>
</SequenceData>
<SequenceData sequenceIDNumber="20">
<INSDSeq>
<INSDSeq_length>15</INSDSeq_length>
<INSDSeq_moltype>AA</INSDSeq_moltype>
<INSDSeq_division>PAT</INSDSeq_division>
<INSDSeq_feature-table>
<INSDFeature>
<INSDFeature_key>source</INSDFeature_key>
<INSDFeature_location>1..15</INSDFeature_location>
<INSDFeature_quals>
<INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>
<INSDQualifier id="q20">

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic construct</INSDQualifier_value>
</INSDQualifier>
</INSDFeature_quals>
</INSDFeature>
</INSDSeq_feature-table>
<INSDSeq_sequence>RASKGVSTSGYSYLH</INSDSeq_sequence>
</INSDSeq>
</SequenceData>
<SequenceData sequenceIDNumber="21">

```
                <INSDSeq>
                        <INSDSeq_length>9</INSDSeq_length>
                        <INSDSeq_moltype>AA</INSDSeq_moltype>
                        <INSDSeq_division>PAT</INSDSeq_division>
                        <INSDSeq_feature-table>
                                <INSDFeature>
                                        <INSDFeature_key>source</INSDFeature_key>
                                        <INSDFeature_location>1..9</INSDFeature_location>
                                        <INSDFeature_quals>
                                                <INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
                                                </INSDQualifier>
                                                <INSDQualifier id="q22">

<INSDQualifier_name>organism</INSDQualifier_name>
                                                        <INSDQualifier_value>synthetic
construct</INSDQualifier_value>
                                                </INSDQualifier>
                                        </INSDFeature_quals>
                                </INSDFeature>
                        </INSDSeq_feature-table>
                        <INSDSeq_sequence>QHSRDLPLT</INSDSeq_sequence>
                </INSDSeq>
        </SequenceData>
        <SequenceData sequenceIDNumber="22">
                <INSDSeq>
                        <INSDSeq_length>7</INSDSeq_length>
                        <INSDSeq_moltype>AA</INSDSeq_moltype>
                        <INSDSeq_division>PAT</INSDSeq_division>
                        <INSDSeq_feature-table>
                                <INSDFeature>
                                        <INSDFeature_key>source</INSDFeature_key>
                                        <INSDFeature_location>1..7</INSDFeature_location>
                                        <INSDFeature_quals>
                                                <INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
                                                </INSDQualifier>
                                                <INSDQualifier id="q21">

<INSDQualifier_name>organism</INSDQualifier_name>
                                                        <INSDQualifier_value>synthetic
construct</INSDQualifier_value>
                                                </INSDQualifier>
```

```
                                        </INSDFeature_quals>
                                </INSDFeature>
                        </INSDSeq_feature-table>
                        <INSDSeq_sequence>LASYLES</INSDSeq_sequence>
                </INSDSeq>
        </SequenceData>
        <SequenceData sequenceIDNumber="23">
                <INSDSeq>
                        <INSDSeq_length>30</INSDSeq_length>
                        <INSDSeq_moltype>AA</INSDSeq_moltype>
                        <INSDSeq_division>PAT</INSDSeq_division>
                        <INSDSeq_feature-table>
                                <INSDFeature>
                                        <INSDFeature_key>source</INSDFeature_key>
                                        <INSDFeature_location>1..30</INSDFeature_location>
                                        <INSDFeature_quals>
                                                <INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
                                                </INSDQualifier>
                                                <INSDQualifier id="q23">

<INSDQualifier_name>organism</INSDQualifier_name>
                                                        <INSDQualifier_value>synthetic
construct</INSDQualifier_value>
                                                </INSDQualifier>
                                        </INSDFeature_quals>
                                </INSDFeature>
                        </INSDSeq_feature-table>

<INSDSeq_sequence>QVQLVQSGVEVKKPGASVKVSCKASGYTFT</INSDSeq_sequence>
                </INSDSeq>
        </SequenceData>
        <SequenceData sequenceIDNumber="24">
                <INSDSeq>
                        <INSDSeq_length>14</INSDSeq_length>
                        <INSDSeq_moltype>AA</INSDSeq_moltype>
                        <INSDSeq_division>PAT</INSDSeq_division>
                        <INSDSeq_feature-table>
                                <INSDFeature>
                                        <INSDFeature_key>source</INSDFeature_key>
                                        <INSDFeature_location>1..14</INSDFeature_location>
                                        <INSDFeature_quals>
                                                <INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>
```

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>
<INSDQualifier id="q24">

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic construct</INSDQualifier_value>
</INSDQualifier>
</INSDFeature_quals>
</INSDFeature>
</INSDSeq_feature-table>
<INSDSeq_sequence>WVRQAPGQGLEWMG</INSDSeq_sequence>
</INSDSeq>
</SequenceData>
<SequenceData sequenceIDNumber="25">
<INSDSeq>
<INSDSeq_length>32</INSDSeq_length>
<INSDSeq_moltype>AA</INSDSeq_moltype>
<INSDSeq_division>PAT</INSDSeq_division>
<INSDSeq_feature-table>
<INSDFeature>
<INSDFeature_key>source</INSDFeature_key>
<INSDFeature_location>1..32</INSDFeature_location>
<INSDFeature_quals>
<INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>
<INSDQualifier id="q25">

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic construct</INSDQualifier_value>
</INSDQualifier>
</INSDFeature_quals>
</INSDFeature>
</INSDSeq_feature-table>

<INSDSeq_sequence>RVTLTTDSSTTTAYMELKSLQFDDTAVYYCAR</INSDSeq_sequence>
</INSDSeq>
</SequenceData>
<SequenceData sequenceIDNumber="26">
<INSDSeq>
<INSDSeq_length>11</INSDSeq_length>
<INSDSeq_moltype>AA</INSDSeq_moltype>
<INSDSeq_division>PAT</INSDSeq_division>
<INSDSeq_feature-table>

<INSDFeature>
<INSDFeature_key>source</INSDFeature_key>
<INSDFeature_location>1..11</INSDFeature_location>
<INSDFeature_quals>
<INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>
<INSDQualifier id="q26">

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic construct</INSDQualifier_value>
</INSDQualifier>
</INSDFeature_quals>
</INSDFeature>
</INSDSeq_feature-table>
<INSDSeq_sequence>WGQGTTVTVSS</INSDSeq_sequence>
</INSDSeq>
</SequenceData>
<SequenceData sequenceIDNumber="27">
<INSDSeq>
<INSDSeq_length>23</INSDSeq_length>
<INSDSeq_moltype>AA</INSDSeq_moltype>
<INSDSeq_division>PAT</INSDSeq_division>
<INSDSeq_feature-table>
<INSDFeature>
<INSDFeature_key>source</INSDFeature_key>
<INSDFeature_location>1..23</INSDFeature_location>
<INSDFeature_quals>
<INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>
<INSDQualifier id="q27">

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic construct</INSDQualifier_value>
</INSDQualifier>
</INSDFeature_quals>
</INSDFeature>
</INSDSeq_feature-table>
<INSDSeq_sequence>EIVLTQSPATLSLSPGERATLSC</INSDSeq_sequence>
</INSDSeq>

```
        </SequenceData>
        <SequenceData sequenceIDNumber="28">
                <INSDSeq>
                        <INSDSeq_length>15</INSDSeq_length>
                        <INSDSeq_moltype>AA</INSDSeq_moltype>
                        <INSDSeq_division>PAT</INSDSeq_division>
                        <INSDSeq_feature-table>
                                <INSDFeature>
                                        <INSDFeature_key>source</INSDFeature_key>
                                        <INSDFeature_location>1..15</INSDFeature_location>
                                        <INSDFeature_quals>
                                                <INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
                                                </INSDQualifier>
                                                <INSDQualifier id="q28">

<INSDQualifier_name>organism</INSDQualifier_name>
                                                        <INSDQualifier_value>synthetic
construct</INSDQualifier_value>
                                                </INSDQualifier>
                                        </INSDFeature_quals>
                                </INSDFeature>
                        </INSDSeq_feature-table>
                        <INSDSeq_sequence>WYQQKPGQAPRLLIY</INSDSeq_sequence>
                </INSDSeq>
        </SequenceData>
        <SequenceData sequenceIDNumber="29">
                <INSDSeq>
                        <INSDSeq_length>32</INSDSeq_length>
                        <INSDSeq_moltype>AA</INSDSeq_moltype>
                        <INSDSeq_division>PAT</INSDSeq_division>
                        <INSDSeq_feature-table>
                                <INSDFeature>
                                        <INSDFeature_key>source</INSDFeature_key>
                                        <INSDFeature_location>1..32</INSDFeature_location>
                                        <INSDFeature_quals>
                                                <INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
                                                </INSDQualifier>
                                                <INSDQualifier id="q29">

<INSDQualifier_name>organism</INSDQualifier_name>
                                                        <INSDQualifier_value>synthetic
```

```
construct</INSDQualifier_value>
                                                    </INSDQualifier>
                                                </INSDFeature_quals>
                                            </INSDFeature>
                                        </INSDSeq_feature-table>

<INSDSeq_sequence>GVPARFSGSGSGTDFTLTISSLEPEDFAVYYC</INSDSeq_sequence>
                    </INSDSeq>
                </SequenceData>
                <SequenceData sequenceIDNumber="30">
                    <INSDSeq>
                        <INSDSeq_length>10</INSDSeq_length>
                        <INSDSeq_moltype>AA</INSDSeq_moltype>
                        <INSDSeq_division>PAT</INSDSeq_division>
                        <INSDSeq_feature-table>
                            <INSDFeature>
                                <INSDFeature_key>source</INSDFeature_key>
                                <INSDFeature_location>1..10</INSDFeature_location>
                                <INSDFeature_quals>
                                    <INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
                                    </INSDQualifier>
                                    <INSDQualifier id="q30">

<INSDQualifier_name>organism</INSDQualifier_name>
                                        <INSDQualifier_value>synthetic
construct</INSDQualifier_value>
                                    </INSDQualifier>
                                </INSDFeature_quals>
                            </INSDFeature>
                        </INSDSeq_feature-table>
                        <INSDSeq_sequence>FGGGTKVEIK</INSDSeq_sequence>
                    </INSDSeq>
                </SequenceData>
                <SequenceData sequenceIDNumber="31">
                    <INSDSeq>
                        <INSDSeq_length>120</INSDSeq_length>
                        <INSDSeq_moltype>AA</INSDSeq_moltype>
                        <INSDSeq_division>PAT</INSDSeq_division>
                        <INSDSeq_feature-table>
                            <INSDFeature>
                                <INSDFeature_key>source</INSDFeature_key>
                                <INSDFeature_location>1..120</INSDFeature_location>
                                <INSDFeature_quals>
                                    <INSDQualifier>
```

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>
<INSDQualifier id="q31">

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic construct</INSDQualifier_value>
</INSDQualifier>
</INSDFeature_quals>
</INSDFeature>
</INSDSeq_feature-table>

<INSDSeq_sequence>QVQLVQSGVEVKKPGASVKVSCKASGYTFTNYYMYWVRQAPGQGLEWMGGINPSNGGTNFNEKFKNRVTLTTDSSTTTAYMELKSLQFDDTAVYYCARRDYRFDMGFDYWGQGTTVTVSS</INSDSeq_sequence>
</INSDSeq>
</SequenceData>
<SequenceData sequenceIDNumber="32">
<INSDSeq>
<INSDSeq_length>111</INSDSeq_length>
<INSDSeq_moltype>AA</INSDSeq_moltype>
<INSDSeq_division>PAT</INSDSeq_division>
<INSDSeq_feature-table>
<INSDFeature>
<INSDFeature_key>source</INSDFeature_key>
<INSDFeature_location>1..111</INSDFeature_location>
<INSDFeature_quals>
<INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>
<INSDQualifier id="q32">

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic construct</INSDQualifier_value>
</INSDQualifier>
</INSDFeature_quals>
</INSDFeature>
</INSDSeq_feature-table>

<INSDSeq_sequence>EIVLTQSPATLSLSPGERATLSCRASKGVSTSGYSYLHWYQQKPGQAPRLLIYLASYLESGVPARFSGSGSGTDFTLTISSLEPEDFAVYYCQHSRDLPLTFGGGTKVEIK</INSDSeq_sequence>
</INSDSeq>
</SequenceData>
<SequenceData sequenceIDNumber="33">

<INSDSeq>
<INSDSeq_length>446</INSDSeq_length>
<INSDSeq_moltype>AA</INSDSeq_moltype>
<INSDSeq_division>PAT</INSDSeq_division>
<INSDSeq_feature-table>
<INSDFeature>
<INSDFeature_key>source</INSDFeature_key>
<INSDFeature_location>1..446</INSDFeature_location>
<INSDFeature_quals>
<INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>
<INSDQualifier id="q33">

<INSDQualifier_name>organism</INSDQualifier_name>
<INSDQualifier_value>synthetic
construct</INSDQualifier_value>
</INSDQualifier>
</INSDFeature_quals>
</INSDFeature>
</INSDSeq_feature-table>

<INSDSeq_sequence>QVQLVQSGVEVKKPGASVKVSCKASGYTFTNYYMYWVRQAPGQGLEWMGGINPSNGGTNFNEKF
KNRVTLTTDSSTTTAYMELKSLQFDDTAVYYCARRDYRFDMGFDYWGQGTTVTVSSASTKGPSVFPLAPCSRSTSESTAALGCL
VKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSGLYSLSSVVTVPSSSLGTKTYTCNVDHKPSNTKVDKRVESKYGPPCPPCPAPE
FLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSQEDPEVQFNWYVDGVEVHNAKTKPREEQFNSTYRVVSVLTVLHQDWLNG
KEYKCKVSNKGLPSSIEKTISKAKGQPREPQVYTLPPSQEEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSD
GSFFLYSRLTVDKSRWQEGNVFSCSVMHEALHNHYTQKSLSLSLG</INSDSeq_sequence>
</INSDSeq>
</SequenceData>
<SequenceData sequenceIDNumber="34">
<INSDSeq>
<INSDSeq_length>218</INSDSeq_length>
<INSDSeq_moltype>AA</INSDSeq_moltype>
<INSDSeq_division>PAT</INSDSeq_division>
<INSDSeq_feature-table>
<INSDFeature>
<INSDFeature_key>source</INSDFeature_key>
<INSDFeature_location>1..218</INSDFeature_location>
<INSDFeature_quals>
<INSDQualifier>

<INSDQualifier_name>mol_type</INSDQualifier_name>

<INSDQualifier_value>protein</INSDQualifier_value>
</INSDQualifier>

<INSDQualifier id="q34">

<INSDQualifier_name>organism</INSDQualifier_name>

<INSDQualifier_value>synthetic construct</INSDQualifier_value>

</INSDQualifier>

</INSDFeature_quals>

</INSDFeature>

</INSDSeq_feature-table>

<INSDSeq_sequence>EIVLTQSPATLSLSPGERATLSCRASKGVSTSGYSYLHWYQQKPGQAPRLLIYLASYLESGVPARFSGSG SGTDFTLTISSLEPEDFAVYYCQHSRDLPLTFGGGTKVEIKRTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDN ALQSGNSQESVTEQDSKDSTYSLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC</INSDSeq_sequence>

</INSDSeq>

</SequenceData>

</ST26SequenceListing>