SEQUENCE LISTING

<110> INSTITUT PASTEUR OF SHANGHAI, CHINESE ACADEMY OF SCIENCES

<120> COMPOSITION AND COMBINED MEDICATION METHOD FOR TREATING ENTEROVIRUS INFECTION

<130> P2016-2164

<150> CN201610067164.5
<151> 2016-01-29

<150> CN201610067175.3
<151> 2016-01-29

<160> 1

<170> PatentIn version 3.5

<210> 1
<211> 462
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<213> Enterovirus

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Gly Glu Ile Gln Trp Val Lys Pro Asn Lys Glu Thr Gly Arg Leu Asn
1 5 10 15

Ile Asn Gly Pro Thr Arg Thr Lys Leu Glu Pro Ser Val Phe His Asp
20 25 30

Ile Phe Glu Gly Asn Lys Glu Pro Ala Val Leu His Ser Lys Asp Pro
35 40 45

Arg Leu Glu Val Asp Phe Glu Gln Ala Leu Phe Ser Lys Tyr Val Gly
50 55 60

Asn Thr Leu Tyr Glu Pro Asp Glu Tyr Ile Lys Glu Ala Ala Leu His
65 70 75 80

Tyr Ala Asn Gln Leu Lys Gln Leu Glu Ile Asn Thr Ser Gln Met Ser
85 90 95

Met Glu Glu Ala Cys Tyr Gly Thr Glu Asn Leu Glu Ala Ile Asp Leu
100 105 110

His Thr Ser Ala Gly Tyr Pro Tyr Ser Ala Leu Gly Ile Lys Lys Arg
115 120 125

Asp Ile Leu Asp Pro Thr Thr Arg Asp Val Ser Lys Met Lys Phe Tyr
130 135 140

Met Asp Lys Tyr Gly Leu Asp Leu Pro Tyr Ser Thr Tyr Val Lys Asp

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145                 150                 155                 160

Glu Leu Arg Ser Ile Asp Lys Ile Lys Lys Gly Lys Ser Arg Leu Ile
                165                 170                 175

Glu Ala Ser Ser Leu Asn Asp Ser Val Tyr Leu Arg Met Thr Phe Gly
            180                 185                 190

His Leu Tyr Glu Ala Phe His Ala Asn Pro Gly Thr Ile Thr Gly Ser
        195                 200                 205

Ala Val Gly Cys Asn Pro Asp Thr Phe Trp Ser Lys Leu Pro Ile Leu
    210                 215                 220

Leu Pro Gly Ser Leu Phe Ala Phe Asp Tyr Ser Gly Tyr Asp Ala Ser
225                 230                 235                 240

Leu Ser Pro Val Trp Phe Arg Ala Leu Glu Leu Val Leu Arg Glu Ile
                245                 250                 255

Gly Tyr Ser Glu Arg Ala Val Ser Leu Ile Glu Gly Ile Asn His Thr
            260                 265                 270

His His Val Tyr Arg Asn Lys Thr Tyr Cys Val Leu Gly Gly Met Pro
        275                 280                 285

Ser Gly Cys Ser Gly Thr Ser Ile Phe Asn Ser Met Ile Asn Asn Ile
    290                 295                 300

Ile Ile Arg Ala Leu Leu Ile Lys Thr Phe Lys Gly Ile Asp Leu Asp
305                 310                 315                 320

Glu Leu Asn Met Val Ala Tyr Gly Asp Asp Val Leu Ala Ser Tyr Pro
                325                 330                 335

Phe Pro Ile Asp Cys Leu Glu Leu Ala Lys Thr Gly Lys Glu Tyr Gly
            340                 345                 350

Leu Thr Met Thr Pro Ala Asp Lys Ser Pro Cys Phe Asn Glu Val Asn
        355                 360                 365

Trp Gly Asn Ala Thr Phe Leu Lys Arg Gly Phe Leu Pro Asp Glu Gln
    370                 375                 380

Phe Pro Phe Leu Ile His Pro Thr Met Pro Met Arg Glu Ile His Glu
385                 390                 395                 400
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Ser Ile Arg Trp Thr Lys Asp Ala Arg Asn Thr Gln Asp His Val Arg
               405                 410                 415

Ser Leu Cys Leu Leu Ala Trp His Asn Gly Lys Gln Glu Tyr Glu Lys
           420                 425                 430

Phe Val Ser Thr Ile Arg Ser Val Pro Val Gly Arg Ala Leu Ala Ile
       435                 440                 445

Pro Asn Tyr Glu Asn Leu Arg Arg Asn Trp Leu Glu Leu Phe
   450                 455                 460
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