;2:0.000,0.000,1.000,0.966;3:0.000,0.035,1.000,0.965;4:0.000,0.035,1.000,0.965;5:0.000,0.035,1.000,0.965;6:0.000,0.035,1.000,0.965;7:0.000,0.035,1.000,0.965;8:0.000,0.035,1.000,0.965;9:0.000,0.035,1.000,0.965;10:0.000,0.035,1.000,0.965;11:0.000,0.035,1.000,0.965;12:0.000,0.035,1.000,0.965;13:0.000,0.035,1.000,0.965;14:0.000,0.035,1.000,0.965;15:0.000,0.035,1.000,0.965;16:0.000,0.035,1.000,0.965;17:0.000,0.035,1.000,0.965;18:0.000,0.035,1.000,0.965;19:0.000,0.035,1.000,0.965;20:0.000,0.035,1.000,0.965;21:0.000,0.035,1.000,0.965;22:0.000,0.035,1.000,0.965;23:0.000,0.035,1.000,0.965;24:0.000,0.035,1.000,0.965;25:0.000,0.035,1.000,0.965;26:0.000,0.035,1.000,0.965;27:0.000,0.035,1.000,0.965;28:0.000,0.035,1.000,0.965;29:0.000,0.035,1.000,0.965;30:0.000,0.035,1.000,0.965;31:0.000,0.035,1.000,0.965;32:0.000,0.035,1.000,0.965;33:0.000,0.035,1.000,0.965;34:0.000,0.035,1.000,0.965;35:0.000,0.035,1.000,0.965;36:0.000,0.035,1.000,0.965;37:0.000,0.035,1.000,0.965;38:0.000,0.035,1.000,0.965;39:0.000,0.035,1.000,0.965;40:0.000,0.035,1.000,0.965;41:0.000,0.035,1.000,0.965;42:0.000,0.035,1.000,0.965;43:0.000,0.035,1.000,0.965;44:0.000,0.035,1.000,0.965;45:0.000,0.035,1.000,0.965;46:0.000,0.035,1.000,0.965;47:0.000,0.035,1.000,0.965;48:0.000,0.035,1.000,0.965;49:0.000,0.035,1.000,0.965;50:0.000,0.035,1.000,0.965;51:0.000,0.035,1.000,0.965;52:0.000,0.035,1.000,0.965;53:0.000,0.035,1.000,0.965;54:0.000,0.035,1.000,0.965;55:0.000,0.035,1.000,0.965;56:0.000,0.035,1.000,0.965;57:0.000,0.035,1.000,0.965;58:0.000,0.035,1.000,0.965;59:0.000,0.035,1.000,0.965;60:0.000,0.035,1.000,0.965;61:0.000,0.035,1.000,0.965;62:0.000,0.035,1.000,0.965;63:0.000,0.035,1.000,0.965;64:0.000,0.035,1.000,0.965;65:0.000,0.035,1.000,0.965;66:0.000,0.035,1.000,0.965;67:0.000,0.035,1.000,0.965;68:0.000,0.035,1.000,0.965;69:0.000,0.035,1.000,0.965;70:0.000,0.035,1.000,0.965;71:0.000,0.035,1.000,0.965;72:0.000,0.035,1.000,0.965;73:0.000,0.035,1.000,0.965;74:0.000,0.035,1.000,0.965;75:0.000,0.035,1.000,0.965;76:0.000,0.035,1.000,0.965;77:0.000,0.035,1.000,0.965;78:0.000,0.035,1.000,0.965;79:0.000,0.035,1.000,0.965;80:0.000,0.035,1.000,0.965;81:0.000,0.035,1.000,0.965;82:0.000,0.035,1.000,0.965;83:0.000,0.035,1.000,0.965;84:0.000,0.035,1.000,0.965;85:0.000,0.035,1.000,0.965;86:0.000,0.035,1.000,0.965;87:0.000,0.035,1.000,0.965;88:0.000,0.035,1.000,0.965;89:0.000,0.035,1.000,0.965;90:0.000,0.035,1.000,0.965;91:0.000,0.035,1.000,0.965;92:0.000,0.035,1.000,0.965;93:0.000,0.035,1.000,0.965;94:0.000,0.035,1.000,0.965;95:0.000,0.035,1.000,0.965;96:0.000,0.035,1.000,0.965;97:0.000,0.035,1.000,0.965;98:0.000,0.035,1.000,0.965;99:0.000,0.035,1.000,0.965;100:0.000,0.035,1.000,0.965;101:0.000,0.035,1.000,0.965;head decoration protein (CDS_0057);anti-repressor Ant (CDS_0067);transposase (CDS_0061);virion structural protein (CDS_0063);virion structural protein (CDS_0066);head decoration protein (CDS_0058);head decoration protein (CDS_0055);Rha-like transcriptional regulator (CDS_0070);DEDDh_I_II_III_IV_V_VI_1 (CDS_0065);Tad2 anti-Thoeris (APIS029) (CDS_0065);transcriptional regulator (CDS_0064);cas12f2_V-F_2 (CDS_0062);CDS_0060 HNH endonuclease (CDS_0068);CDS_0048;CDS_0049;CDS_0050;CDS_0056;CDS_0057;CDS_0059;CDS_0060;CDS_0061;CDS_0063;CDS_0064;CDS_0069;DNA methyltransferase (CDS_0074);Rha-like transcriptional regulator (CDS_0065);virion structural protein (CDS_0068);virion structural protein (CDS_0069);cas12f2_V-F_2 (CDS_0076);head decoration protein (CDS_0061);head decoration protein (CDS_0059);head decoration protein (CDS_0060);DNA, RNA and nucleotide metabolism;head and packaging;integration and excision;other;transcription regulation;unknown function;Defence;Anti-defence;10000 nt" />