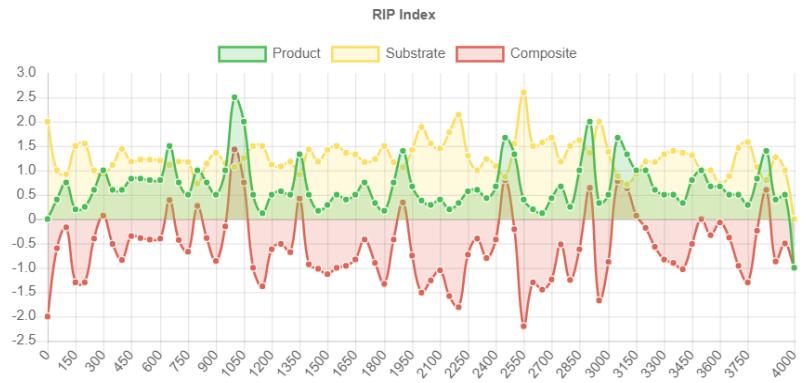
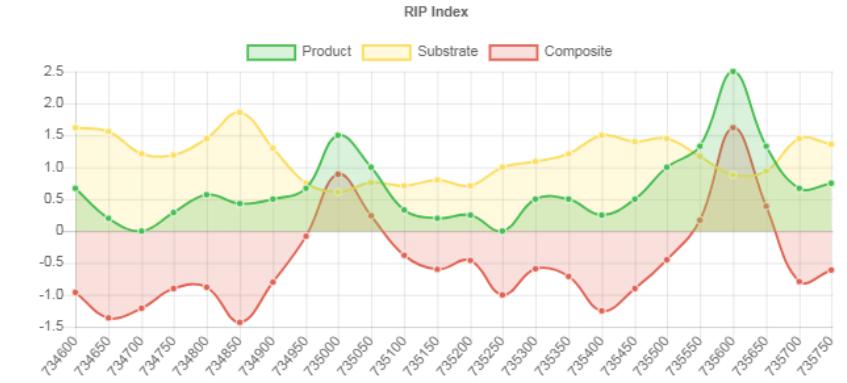
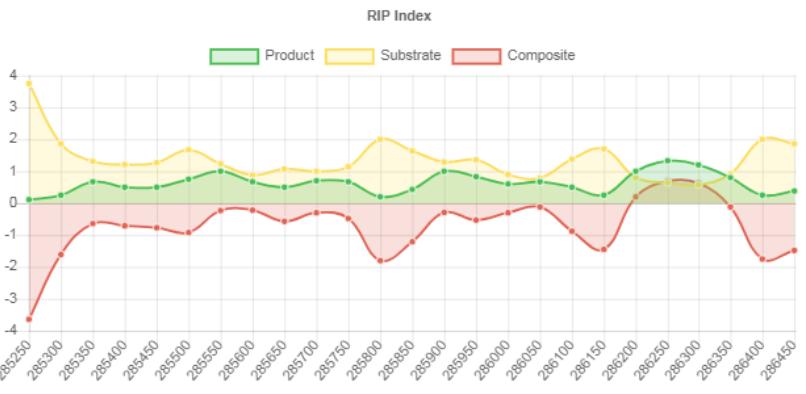
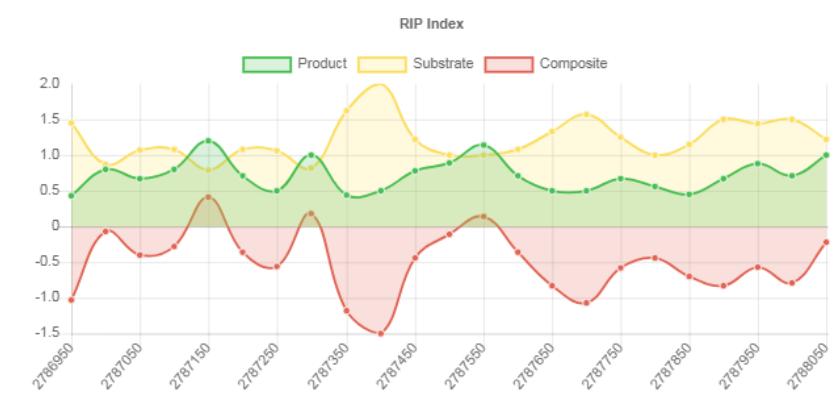
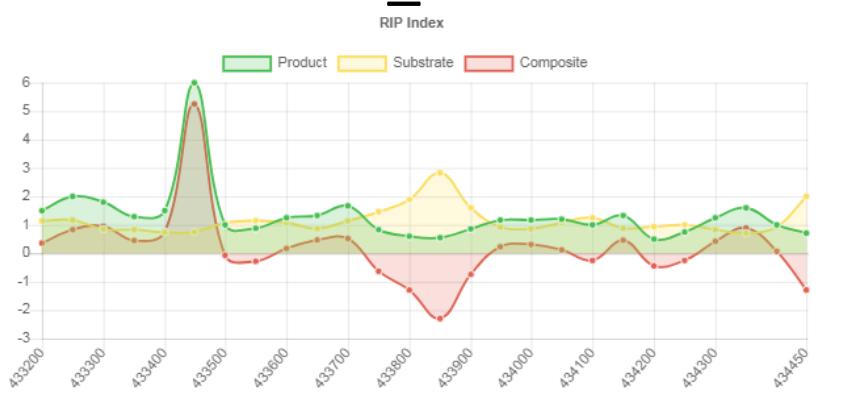
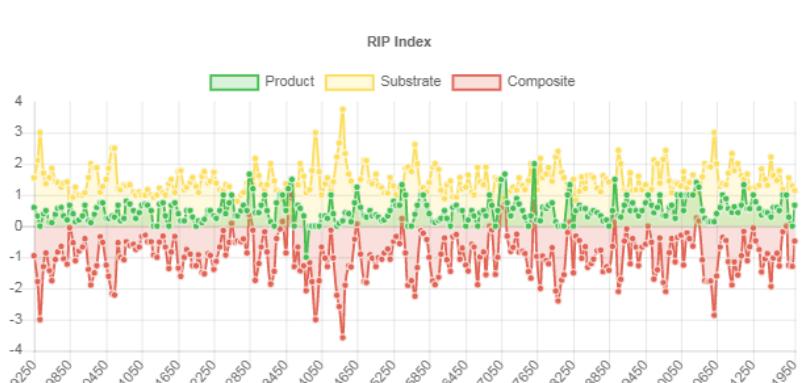
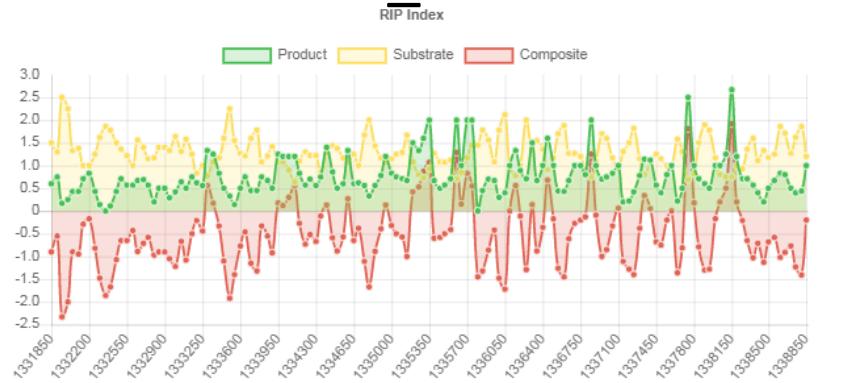
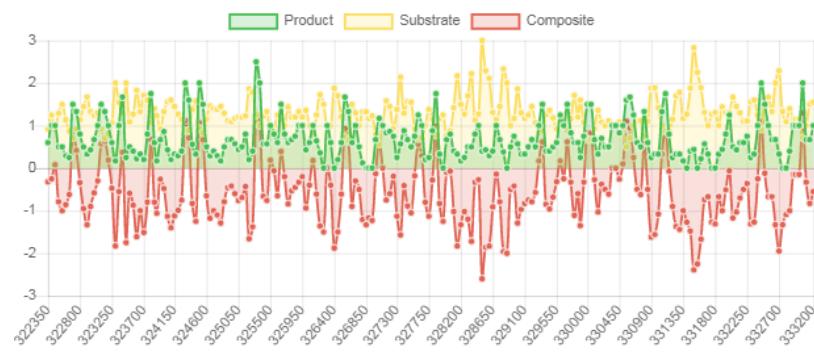


Figure S2: RIP analyses of effector gene dataset: Line graphs depicting the changes in RIP index values (y-axis) across the length of the gene sequences (Table S4). Changes in RIP composite index value (red), RIP substrate index values (yellow) and RIP product index values (green) are indicated. RIP composite values above 0 indicate RIP. RIP substrate index values above 1.1 and RIP substrate index values below 0.75 are suggestive of RIP. The results for each effector gene set are illustrated in Figure S2A to Figure S2E, for FSP34, FSOR, FFRA, CMWF567, and UG27, respectively.

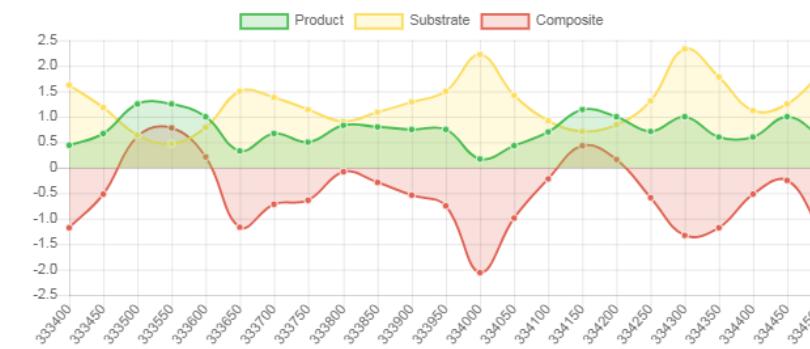
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B**FSOR****g3.48**

RIP Index

**g15.61**

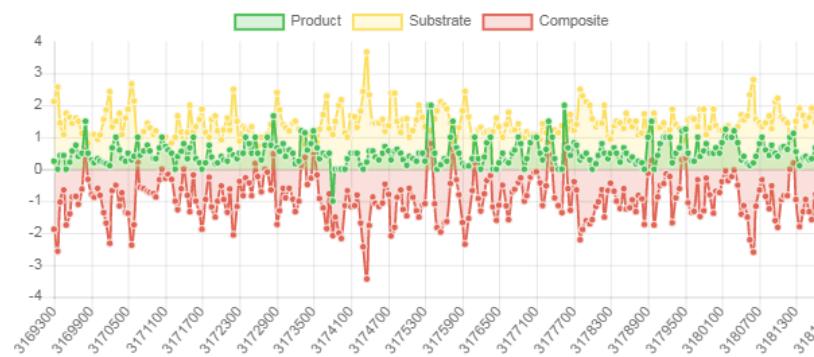
RIP Index

**g39.61**

RIP Index

**g31.74**

RIP Index

**g16.33**

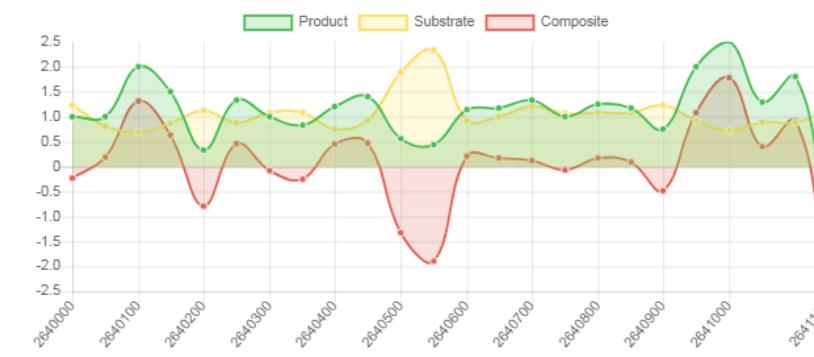
RIP Index

**g2.102**

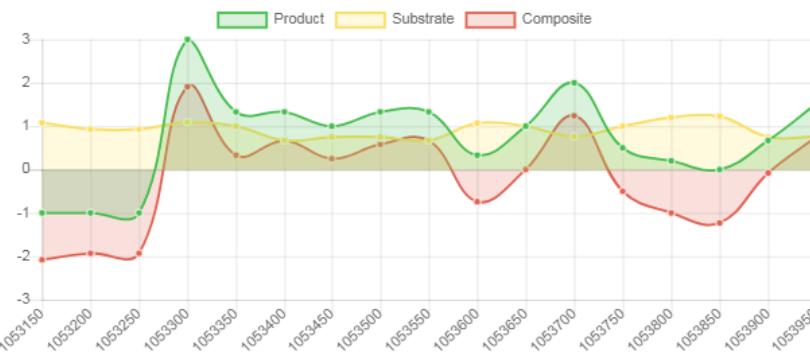
RIP Index

**g26.94**

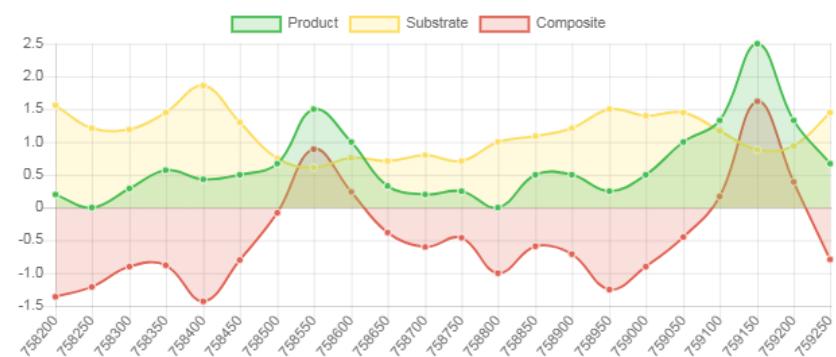
RIP Index

**g10.52**

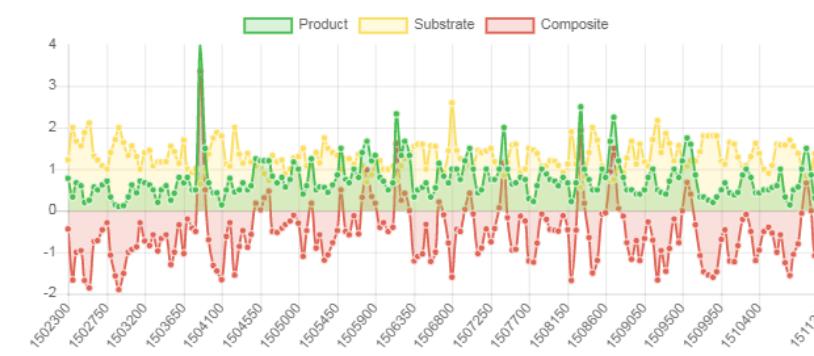
RIP Index

**g7.37**

RIP Index

**g15.66**

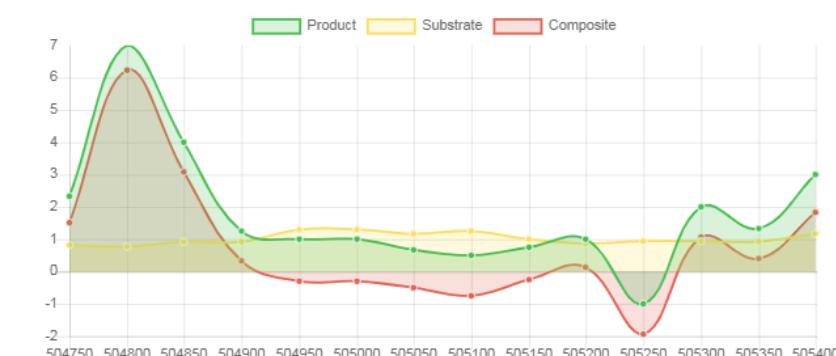
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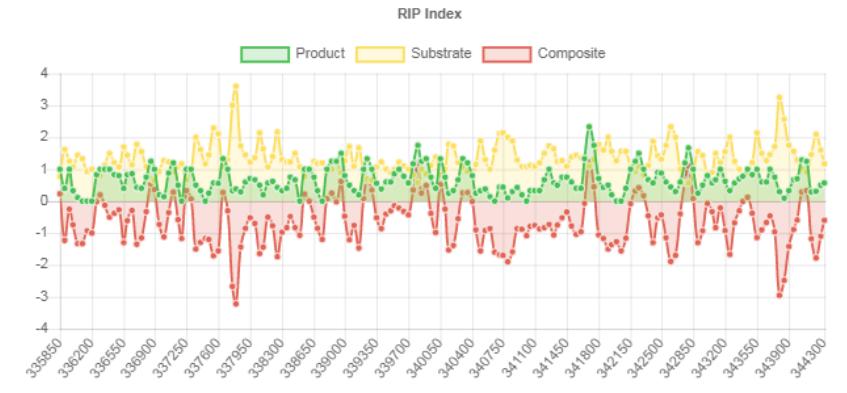
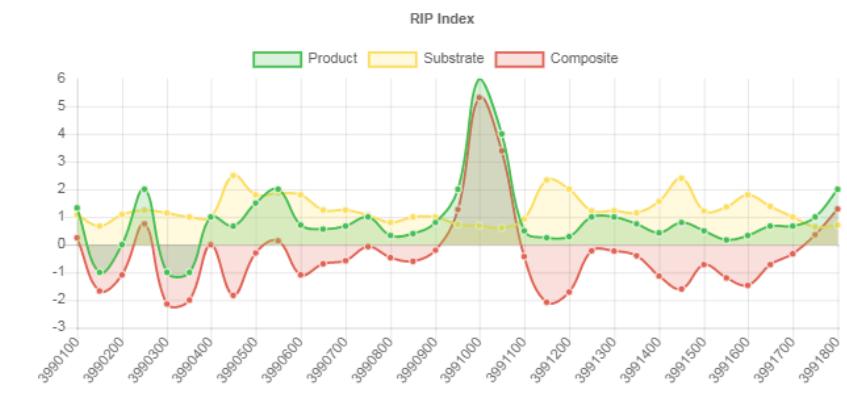
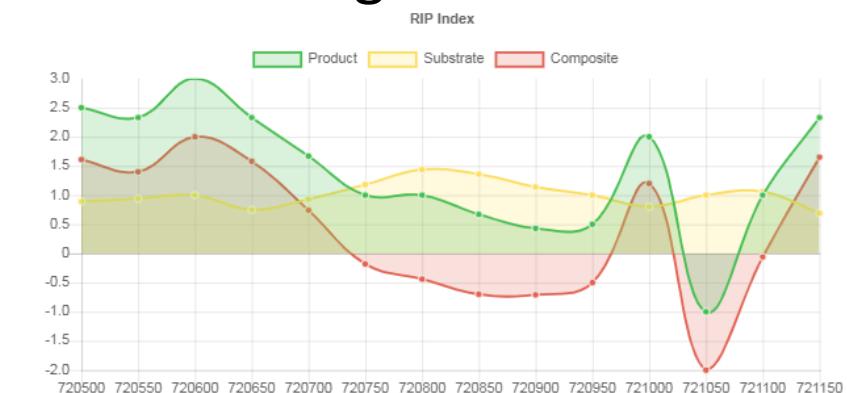
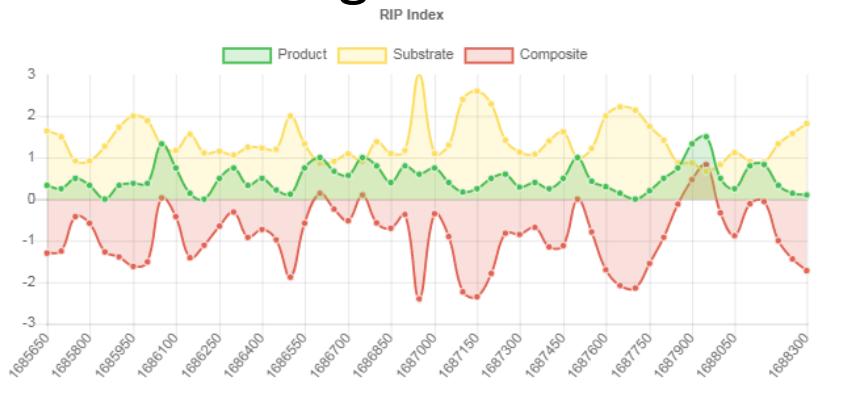
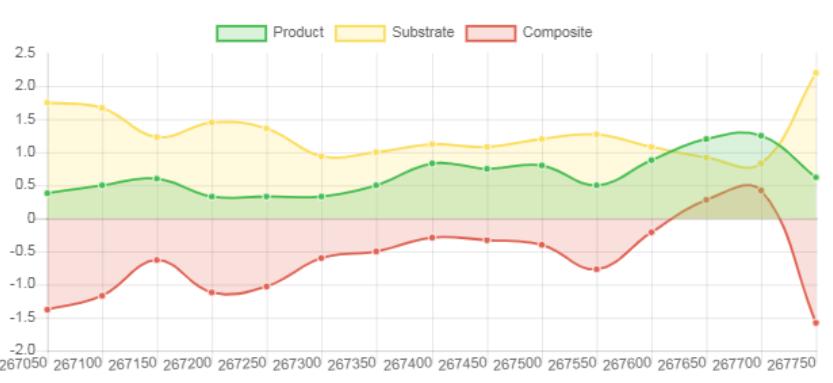
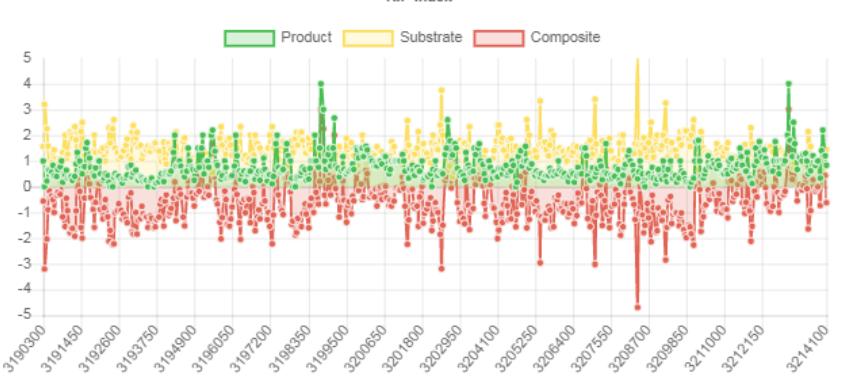
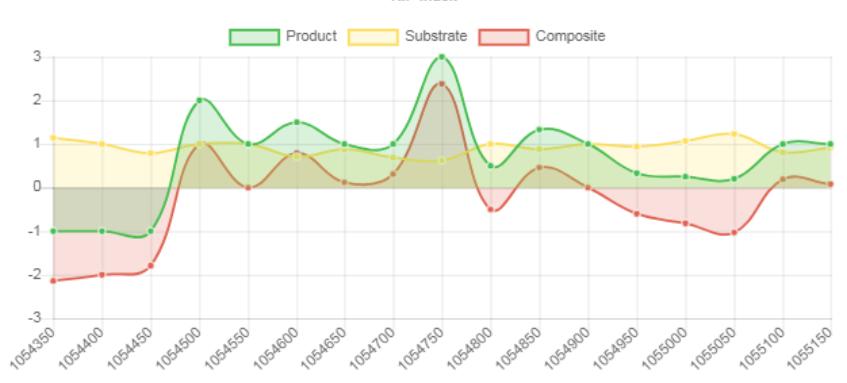
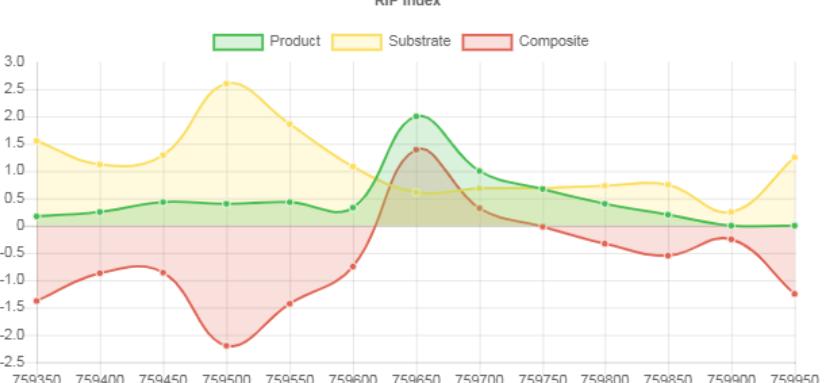
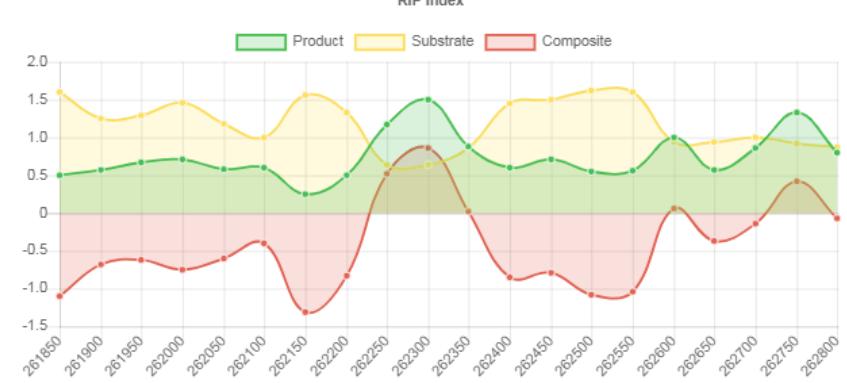
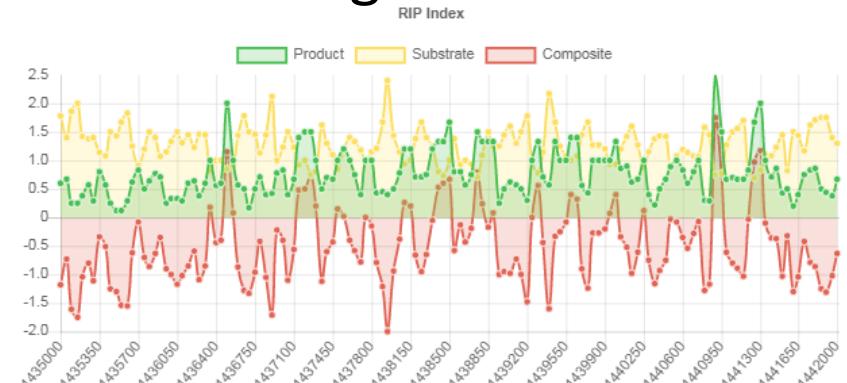
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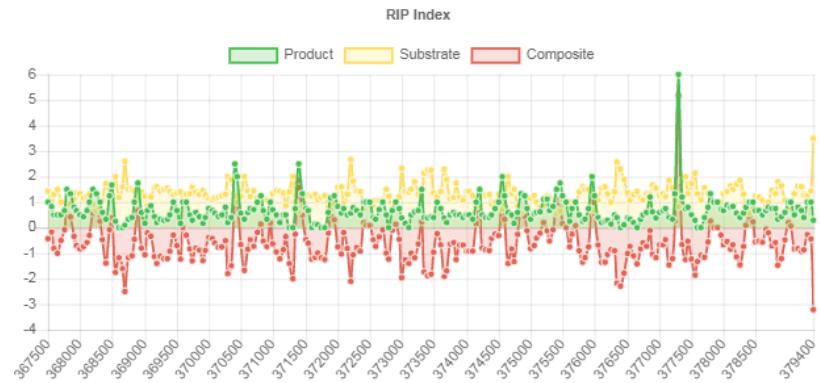
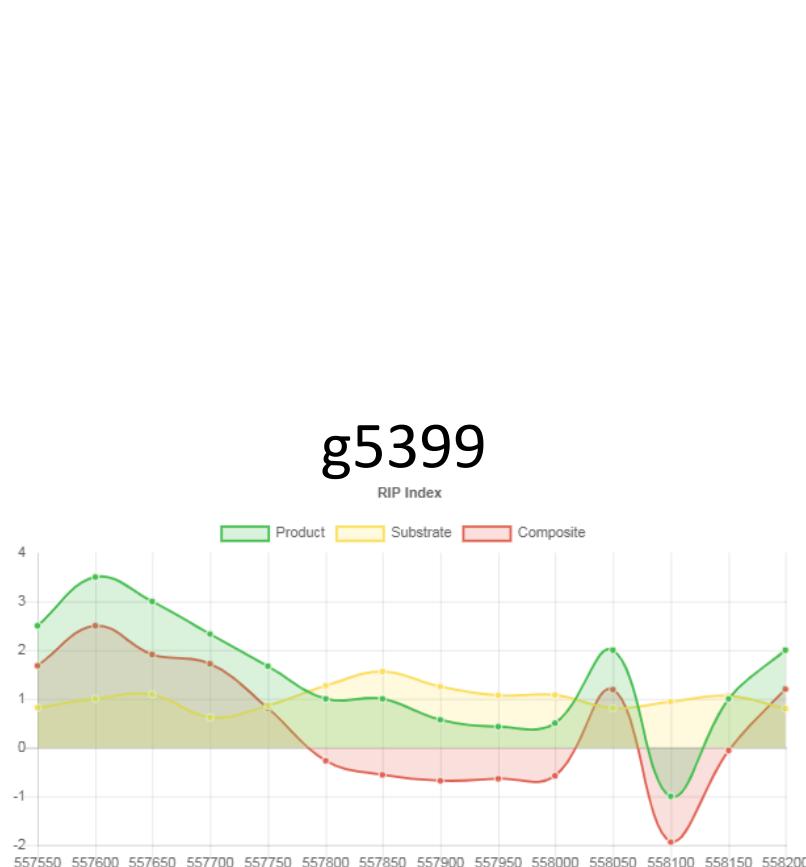
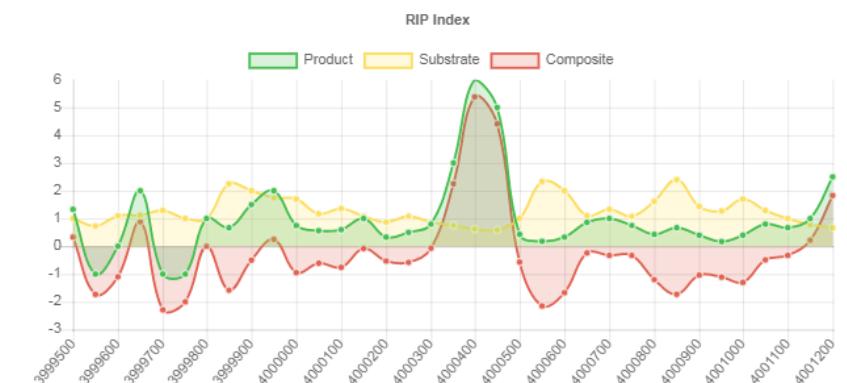
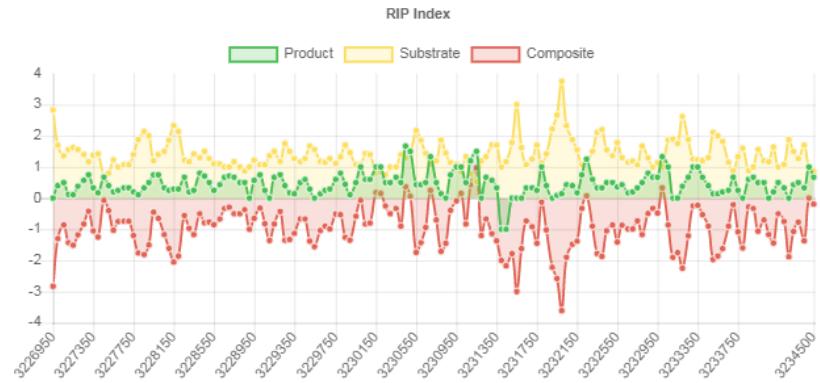
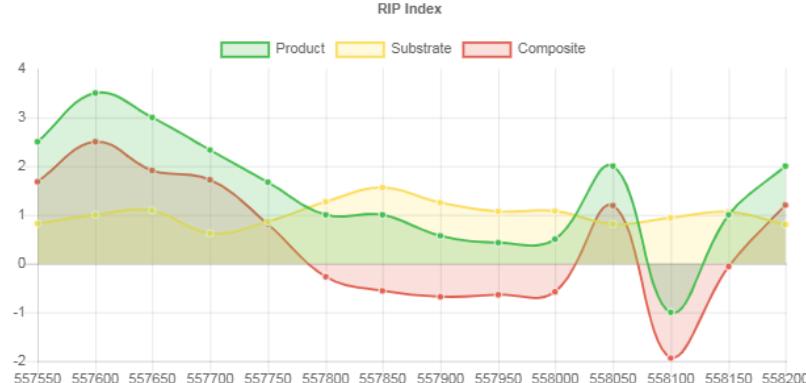
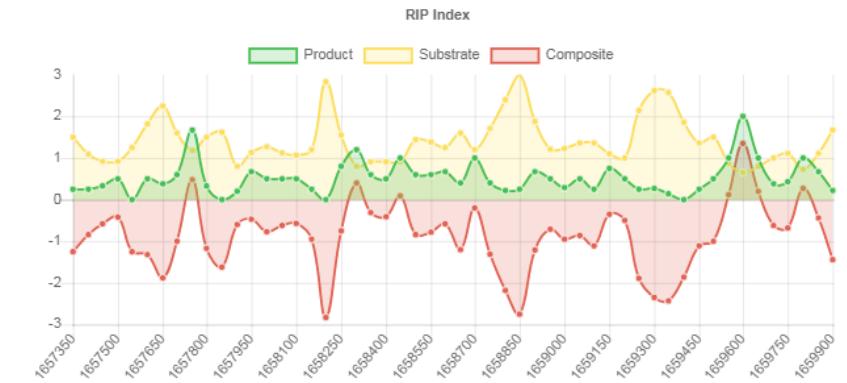
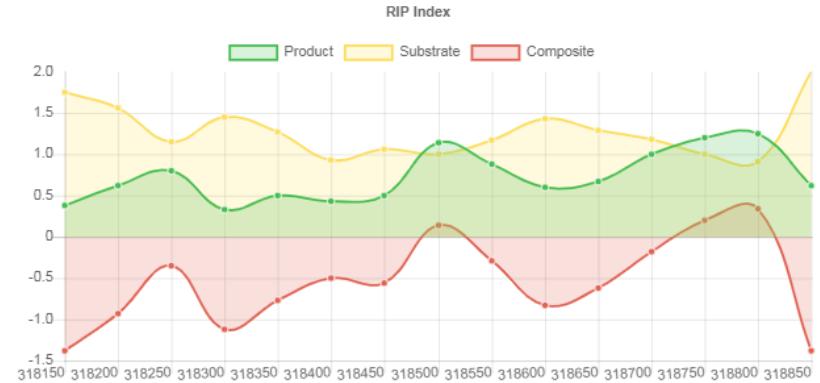
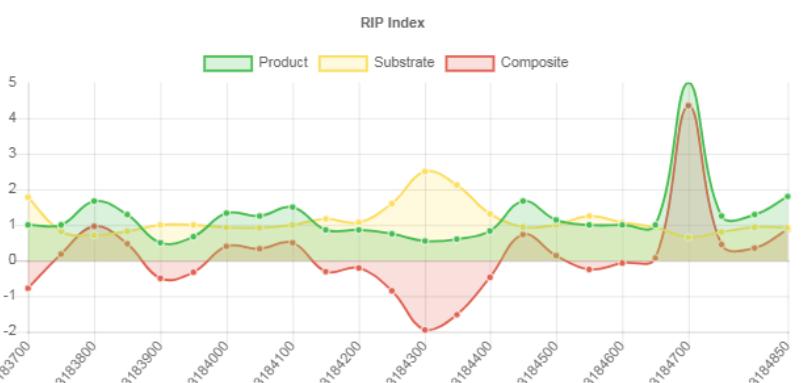
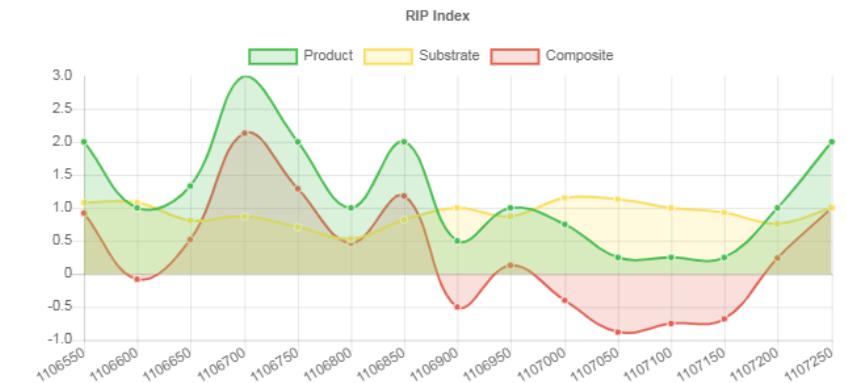
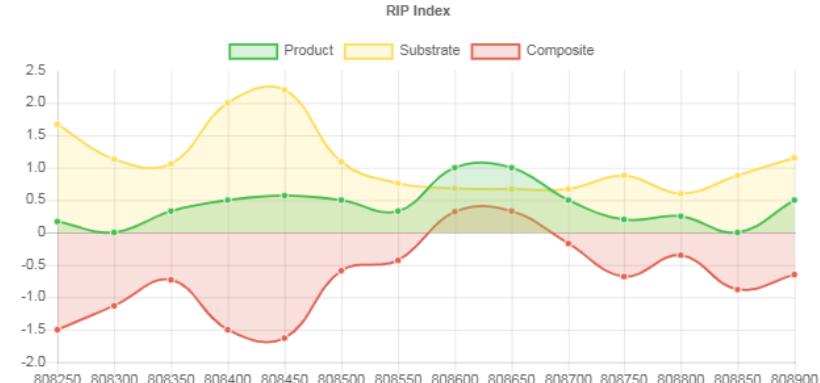
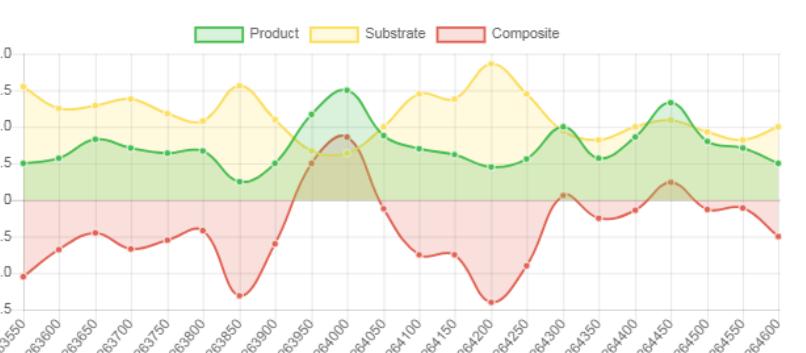
RIP Index

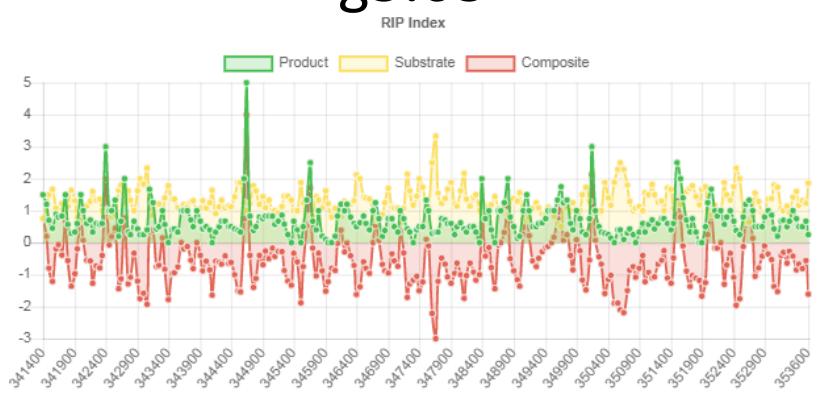
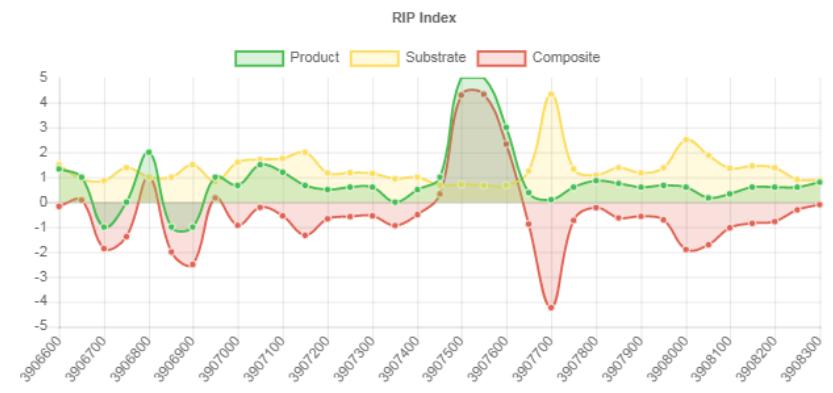
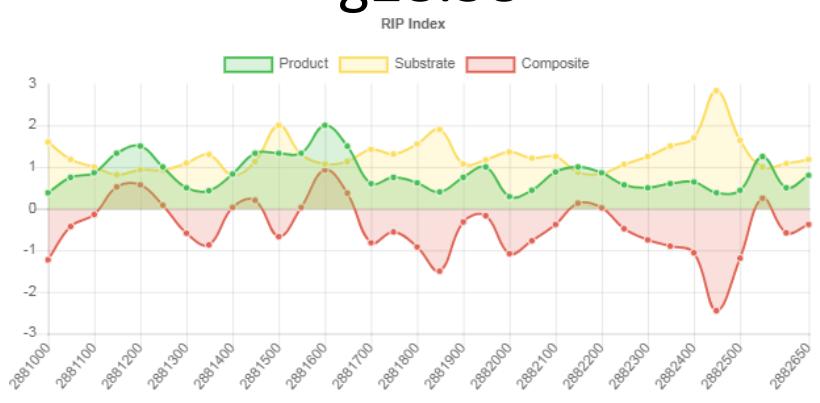
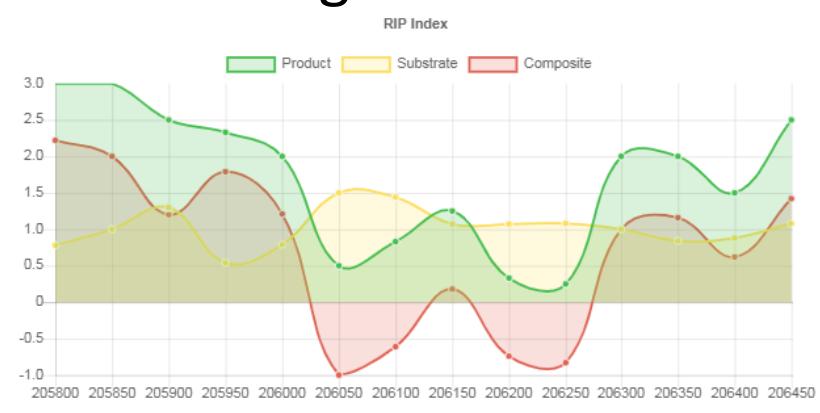
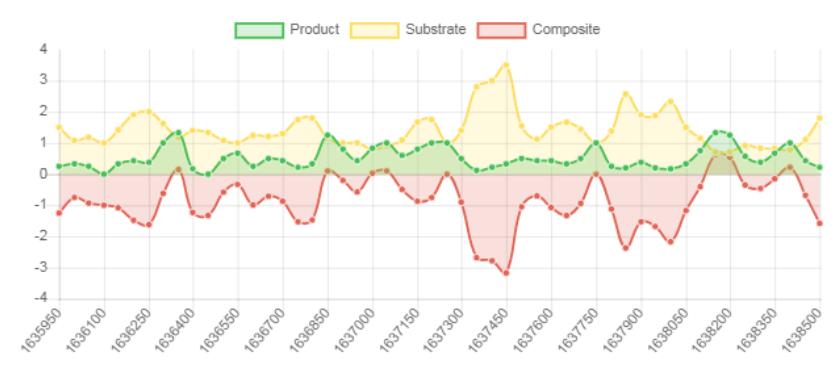
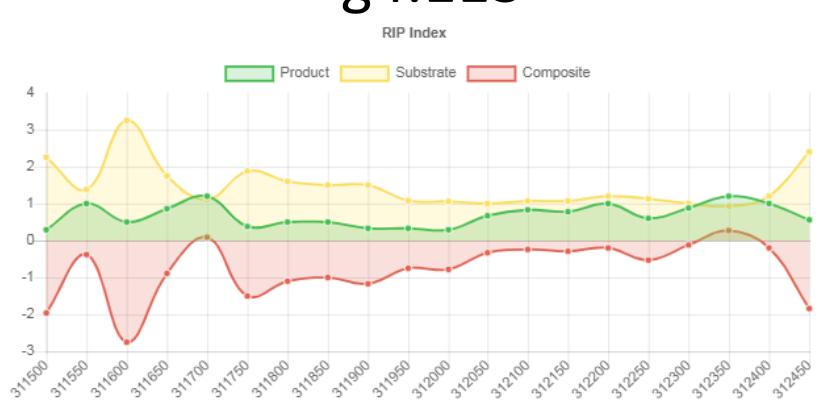
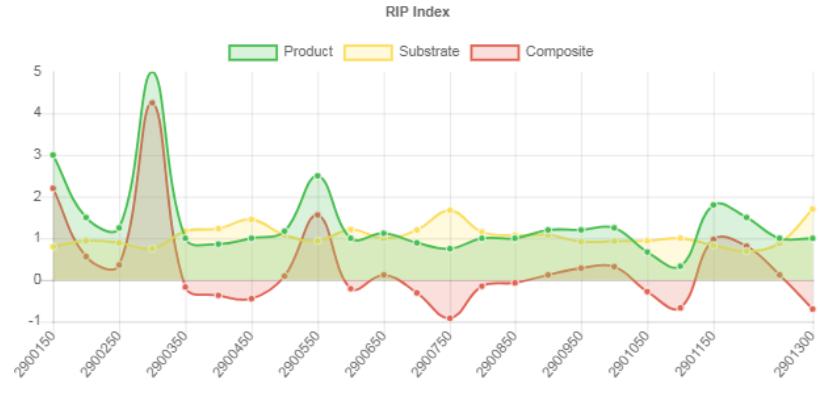
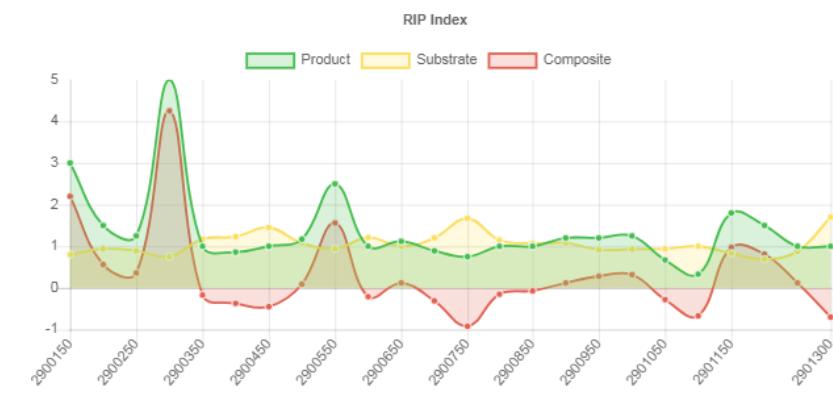
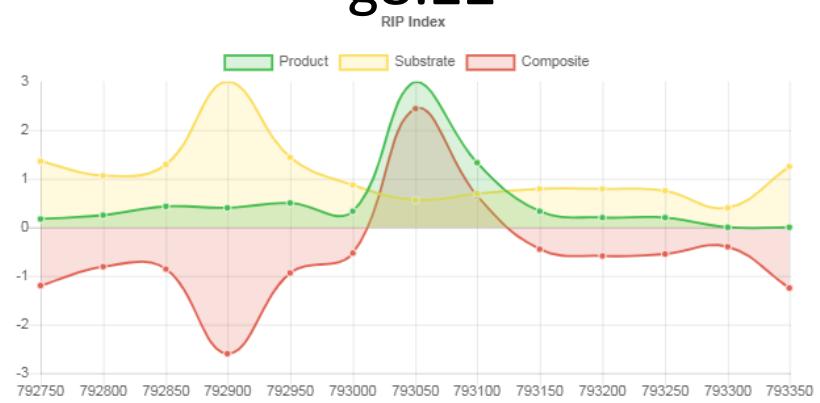
**g5.67**

RIP Index



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E**UG27****g5.63****g39.51****g28.58****g4.107****g16.12****g4.113****g29.43****g10.86****g8.22****g4.9****g14.70**