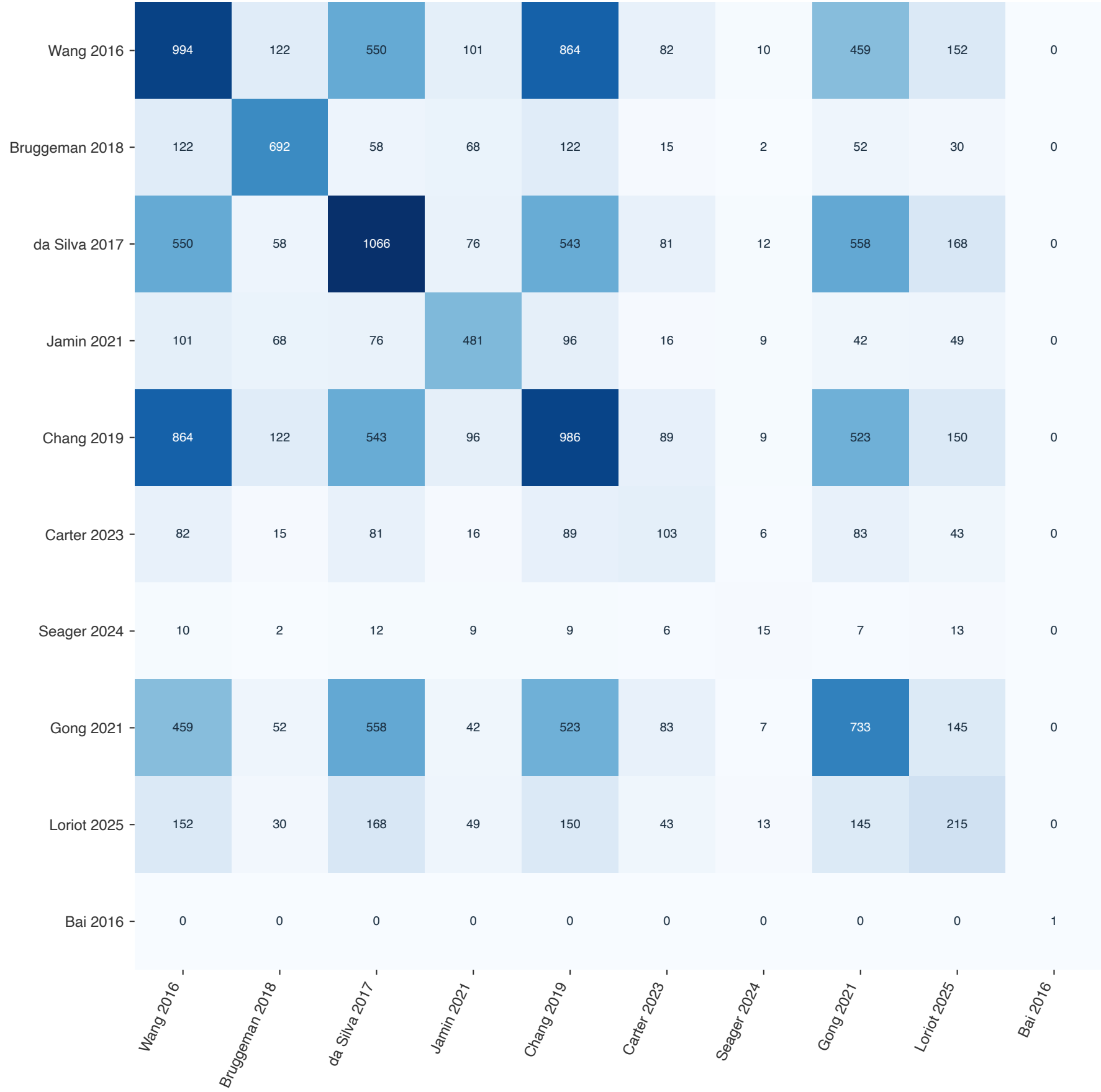


Complete coding candidate pool



Retained by current default rules



Cells count unique shared genes; diagonals are source totals. Nested and overlapping sources must not be summed.