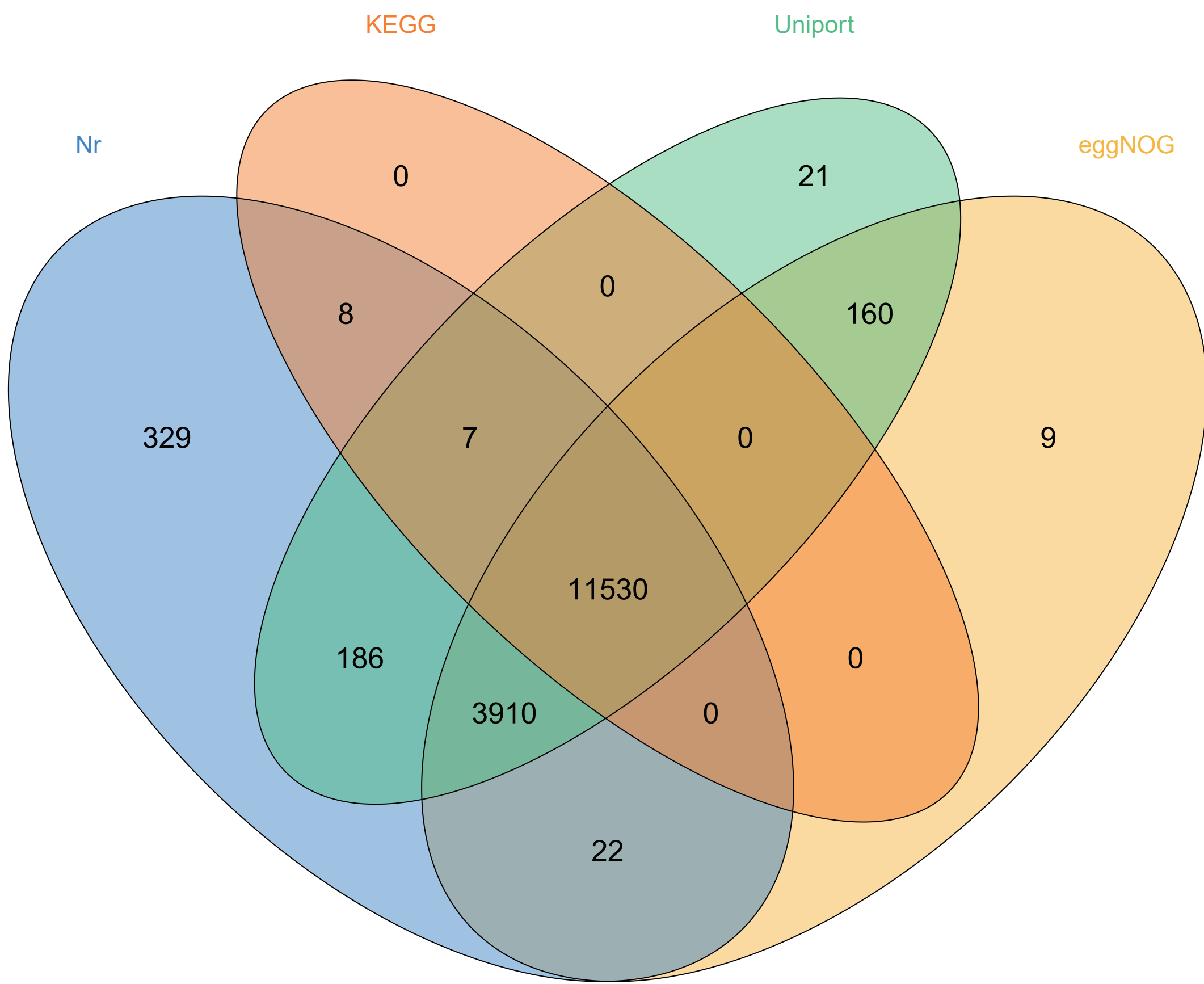




Additional file 5: Fig. S2. Functional annotation of *S. littoralis* predicted genes using 4 databases