



Additional Figure 2. Protein Sequence alignment for Rp-Toll paralogs. Predicted protein sequences for Rp-Toll-2 (Rp-Toll) are aligned with *R. prolixus* Toll paralogs. Light grey to dark grey shading denotes increase in sequence similarity. Region underlined in grey corresponds to the epitope recognized by the anti-Toll antiserum d300 (aa 108-455), and to IL-1-related (aa 1051-1302) region.