

**nanor
orthologs**



Figure S3. **Nanor-like genes are not included in large conserved syntenic blocks.** Phylogenetic relationships between genes located in the neighborhood of zebrafish *nanor* and *nanorb* were studied using genomic and direct blast analysis. Well supported (co)orthology between flanking markers is denoted by red, orange and light blue. Genes represented in pink were significantly similar to medaka ENSORLG00020009526 or to rivulus ENSKAMAG00000012530 (tblastn e values < 1e-10) but did not stand in Ensembl as their true orthologs.