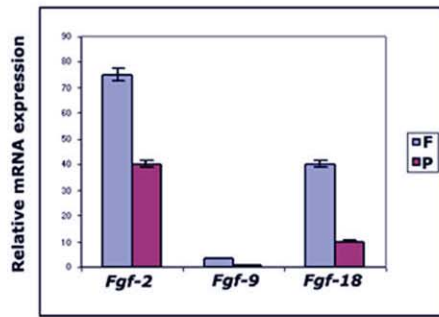
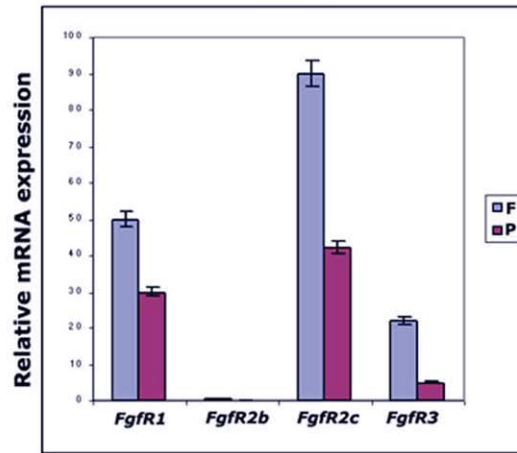


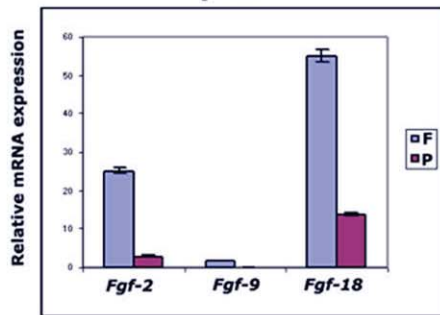
E17.5



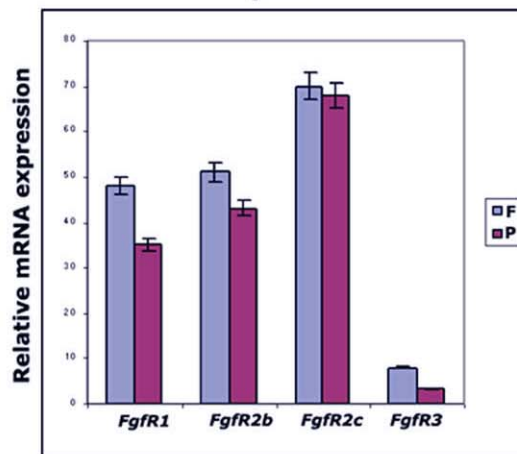
E17.5



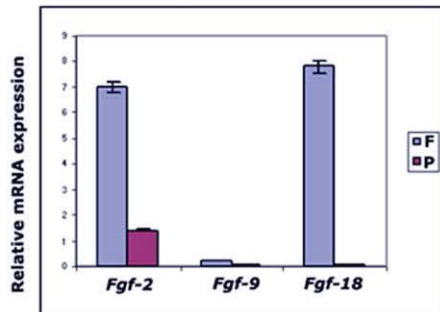
pN1



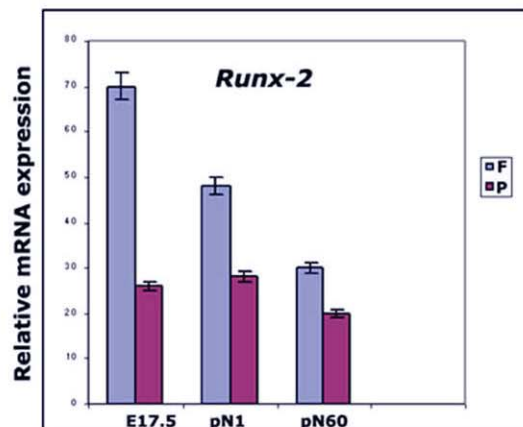
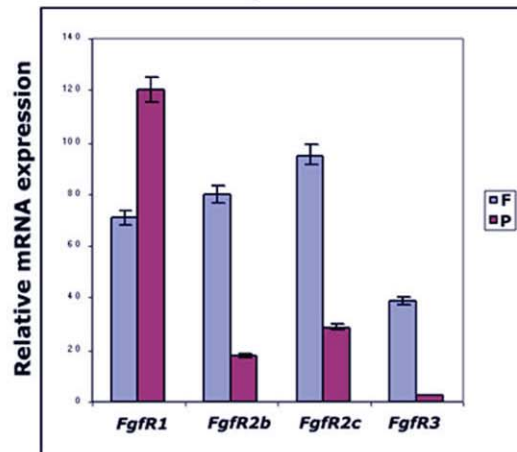
pN1



pN60



pN60



***p value <0.02**

Supplemental Figure 1. Quantitative analysis of *Fgf* ligands and *Fgf* receptors performed by real-time PCR. A standard curve method of quantitation was used to calculate expression of target genes relative to the house keeping gene *Gapdh*. The data correlate with the above results obtained by semiquantitative RT-PCR showing the highest expression level of both *Fgf* ligands and *Fgf* receptors in frontal bone of E17.5, pN1 and pN60. The results are presented as the mean \pm SE of two independent experiments. Statistical differences between the means are examined by Student's test. A. *P* value <0.05 was considered statistically significant.