



Extended Data Figure 6 | Lineage fate of myeloid progenitors. **a**, Two M2/HSB/Tn mice were induced and chased for one week, and then myeloid progenitors (GMP, MEP, and CMP) and Lin⁺ cells were isolated from bone marrow and their transposon tag content was analysed. Chart is a binary representation of all Lin⁺ tags overlapping with any myeloid progenitor tag ranked by lineage. **b**, Quantification of relative lineage

contribution of GMPs, MEPs, and CMPs as a fraction of lineage-specific/total lineage-overlapping clones for each MyP subset. Values are mean of the two analysed mice. **c**, An additional M2/HSB/Tn mouse was induced and chased for three weeks, and then processed as in **a**. **d**, Quantification of relative lineage contribution of GMPs, MEPs, and CMPs at three weeks after labelling.