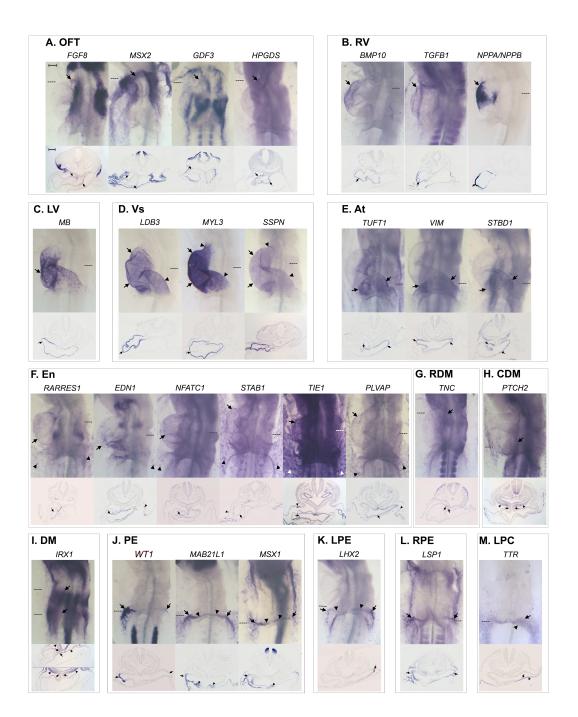


<u>Supplementary figure 1.</u> Hierarchical clustering of RNAseq readings for 132 genes tested by mRNA *in situ* hybridisation. Enriched RNAs are displayed in red, depleted RNA in blue. LV, prospective left ventricle; RV, prospective right ventricle; RDM, rostral dorsal mesocardium; OFT, outflow tract; SV, sinus venosus; LPC, left pacemaker; CDM, caudal dorsal mesocardium; LPE, left proepicardium; RPE, right proepicardium; left-RPC, left side corresponding to the position of the RPC; RPC, right pacemaker.



Supplementary figure 2. A further selection of regionalised heart markers. **A**. *FGF8*, *PRSS23*, *MSX2*, *GDF3/cVg1* and *HPGDS* are detected in the OFT (arrows). **B**. *BMP10* and *TGFBI* expression is restricted to the outer curvature of the prospective RV (arrows). *NPPA/NPPB* expression is observed in the RV (arrow), but also in the inner and outer curvatures of the prospective ventricle. **C**. *MB* is detected in the prospective LV. **D**. *LDB3*, *MYL3* and *SSPN* expression extends beyond the prospective ventricular region (arrows) into the atrial and OFT regions (arrowheads). **E**. *TUFT1*, *VIM* and *STBD1* localise to the atria (arrows). **F**. *RARRES1*, *EDN1*, *NFATC1*, *STAB1*, *TIE1* and *PLVAP* are detected in the endocardium (arrows) and endothelial cells (arrowheads); **G**. *TNC* expression is faint but detectable in the RDM (arrows). **H**. *PTCH2* is weakly detected in the CDM (arrows) and expression is continuous with the splanchnic mesoderm (arrowheads). **I**. *IRX1* in both the rostral and caudal DM (arrows) and the splanchnic mesoderm (arrowheads). **J**. *WT1* transcripts are detected in both the right and the left PE (arrows), as are *MAB21L1*

and MSX1 (arrows) and also in ventro-medial cells (arrowheads). **K**. LPE marker LHX2 is detected in both PE (arrows) and ventro-medial cells (arrowheads). **L**. RPE marker LSP1 is detected in both PEs (arrows). **M**. TTR, a putative marker for the early LPC (arrows), is also expressed in the endoderm (arrowheads). At. Atria; En. Endocardium. Scale bars 100µm.



Supplementary figure 3. Markers with less regional specificity. **A**. *ACTA2*, *RASSF5*, *OLFML2B*, *SPRY2* and *PKIG* transcripts are enriched in the OFT (arrows), but their

expression is not restricted to this region. B. FBXL22 transcripts are enriched (arrows) but not restricted to the prospective RV. AMY2A expression is strong throughout the heart. C. PACAP/ADCYAP1 expression is not restricted to the prospective LV. HS6ST3 expression is faint and PLBD1 and ZNF106 expression is strong throughout the heart. **D**. LOX expression is faint in both the prospective right and left ventricles. PLPPR1 and VCAM1 expression is faint, whereas, CMYA5, EXFABP, MYL1, SORBS2, TNNI2, KCNH6 and LYPD1 are strong in the heart and/or embryo. E. GJD2, FAM3B and UGT1A1 expression is faint in the atrial region (arrows). BORCS8 expression is strong within the embryo and atrial region (arrows). F. Endocardial markers FOXO1, HEG1, POSTN, TAL1 and IQSEC1 are not detected. G. PKDCCB, LUM, BMP4, LOC417741, EBF3 and KAZALD1 expression in the RDM (arrows) is continuous with the splanchnic mesoderm. KAZALD1 and EBF3 are also in the myocardium (arrowheads). PROM1, CCDC3, SIX1 and SLIT2 are faint or not detected in the RDM. H. Expression of CDM markers FOXF2, APCDD1, STRA6 and HOXA2 are faint (arrows). STRA6 is also detected in the endocardium (arrowhead). MYCN, HOXA1, LINGO1 and NGFR are not detected in the CDM. I. RDH10 and SALL4 expression in the DM (arrows) is continuous with the splanchnic mesoderm and the myocardium (arrowheads). LIX1, IRX3, and ER81/ETV1 are not detected in the DM. J. Faint expression of GAL and FGFR2 is detected in both right and left PE (arrows), and GAL is detected in ventro-medial cells (arrowheads). HBZ, HBM and HBAA are not restricted to the PE. K. LPE marker ANXA2 is detected in both PE (arrows). RBP4A, APOC3, HOXB5, LMO2, PAMR1 and HOXB4 are either faint or not expressed. L. GEM, OPRM1 and AGPAT2 are strongly expressed throughout the heart and are not specific to the RPE. M. GUCA2A transcripts are not detected in the LPC region. **N**. CENPW expression is strong throughout the heart and not localised to the RPC region. **O**. JAG1 and RAI14 are detected on both the left (arrow) and right (arrowhead) and is not restricted to the Left-RPC. P. Putative PC marker FGB is expressed in the AIP endoderm and not detected in either the early LPC or RPC. Q. MSI2 and TNR are not restricted to the SV region. Scale bars 100µm.