

Row 3 – H Header row (example)

H,SocAnimId,CalvingYear,Sex,Owned,ced_ebv,ced_acc,ced_perc,ced_prog,cem_ebv,cem_acc,cem_perc,cem_prog,gl_ebv,gl_acc,gl_perc,gl_prog,bw_ebv,bw_acc,bw_perc,bw_prog,ww_ebv,ww_acc,ww_perc,ww_prog,yw_ebv,yw_acc,yw_perc,yw_prog,fw_ebv,fw_acc,fw_perc,fw_prog,mcw_ebv,mcw_acc,mcw_perc,mcw_prog,milk_ebv,milk_acc,milk_perc,milk_prog,scrotal_ebv,scrotal_acc,scrotal_perc,scrotal_prog,dc_ebv,dc_acc,dc_perc,dc_prog,carc_wt_ebv,carc_wt_acc,carc_wt_perc,carc_wt_prog,carc_ema_ebv,carc_ema_acc,carc_ema_perc,scan_ema_prog,carc_ema_prog,carc_rib_fat_ebv,carc_rib_fat_acc,carc_rib_fat_perc,scan_rib_fat_prog,carc_rib_fat_prog,carc_rump_fat_ebv,carc_rump_fat_acc,carc_rump_fat_perc,scan_rump_fat_prog,carc_rump_fat_prog,carc_rby_ebv,carc_rby_acc,carc_rby_perc,carc_rby_prog,carc_imf_ebv,carc_imf_acc,carc_imf_perc,scan_imf_prog,carc_imf_prog,nfi_f_ebv,nfi_f_acc,nfi_f_perc,nfi_f_prog,docility_ebv,docility_acc,docility_perc,docility_prog,struct_fa_ebv,struct_fa_acc,struct_fa_perc,struct_fa_prog,struct_fc_ebv,struct_fc_acc,struct_fc_perc,struct_fc_prog,struct_ra_ebv,struct_ra_acc,struct_ra_perc,struct_ra_prog,struct_rh_ebv,struct_rh_acc,struct_rh_perc,struct_rh_prog,struct_rs_ebv,struct_rs_acc,struct_rs_perc,struct_rs_prog,bobj_1,bobj_1_perc,bobj_2,bobj_2_perc,bobj_3,bobj_3_perc,bobj_4,bobj_4_perc,NumHerds,ProgAnalysed,GenomicProg,in_dam_inventory,age_at_first_calf,calv_interval,trait_indicators

Where:

- SocAnimID is a unique alphanumeric identifier assigned to the animal by Angus Australia
- Calving Year is the 4 digit year of birth
- Sex is a character abbreviation of the animal sex (M,F,S)
- Owned is 1 if the animal is currently in the submitting member's Herd, or 0 if not (ie it is in the pedigree of an animal in the Herd only)
- This is followed by a string of the trait-codes for the traits and selection indices which are included in the file. Each trait has an EBV, accuracy, percentile band and progeny count combination except :
 - Accuracies are not supplied for selection indices
 - Progeny counts are not supplied for the selection indices
- NumHerds is the number of herds in which this animal has performance recorded progeny
- ProgAnalysed is the number of progeny of this animal that had performance information analysed
- GenomicProg is the number of progeny of this animal that have genomic information
- In_dam_inventory is 'Y' if the animal is in the dam inventory, 'N' otherwise
- Age_at_first_calf is the age in years (to 1 decimal place) of a cow at her first calf birth date. Appears as '-' if not relevant.
- Calv_interval
 - '-' if not relevant
 - 'dnr' if donor dam
 - Otherwise, the average number of days between calves, for a cow
- Trait_indicators is a representation of which traits have been observed. This column is enclosed in double-quotes as it contains commas

- PERCENTILE-GROUPnn is the (top) nn% percentile value for the current calves.
 - Percentile rankings are 01, 05, 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95 and 99.
 - The Calving Year reflects the Year for the rankings.
 - Example:
 - D,PERCENTILE-GROUP35,2017,,,,+1.3,,,,+1.3,,,,-4.8,,,,+3.7,,,,+48,,,,+86,,,,+113,,,,+100,,,,+17,,,,+2.1,,,,-5.0,,,,+64,,,,+5.9,,,,+0.5,,,,+0.4,,,,+0.8,,,,+2.0,,,,+0.03,,,,+8,,,,+4,,,,+5,,,,+2,,,,+0.6,,,,+0.1,,,,+121,,+113,,+130,,+117,,,,,