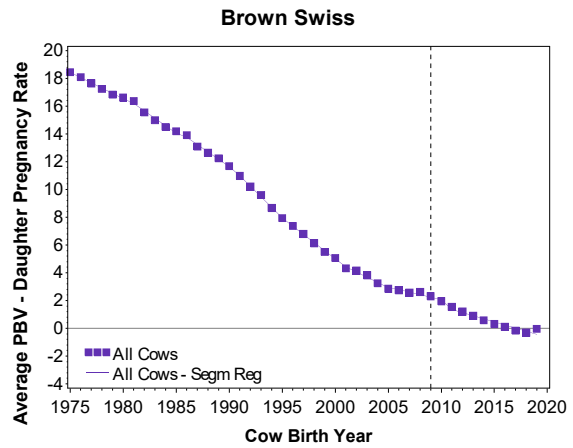
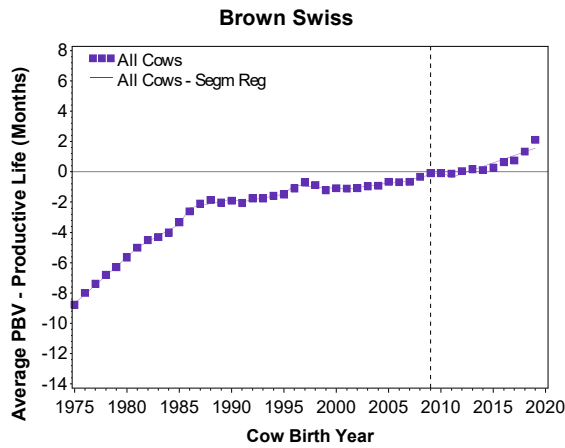
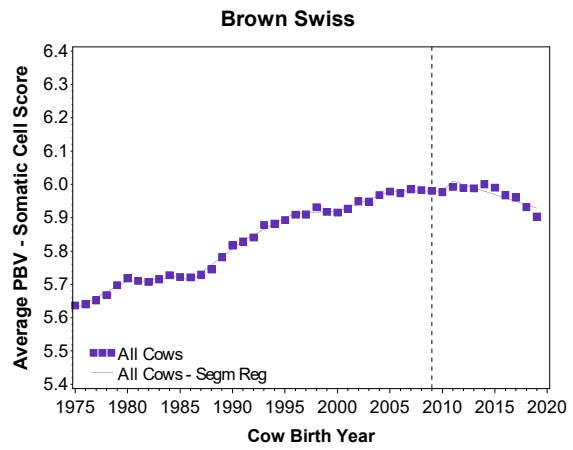
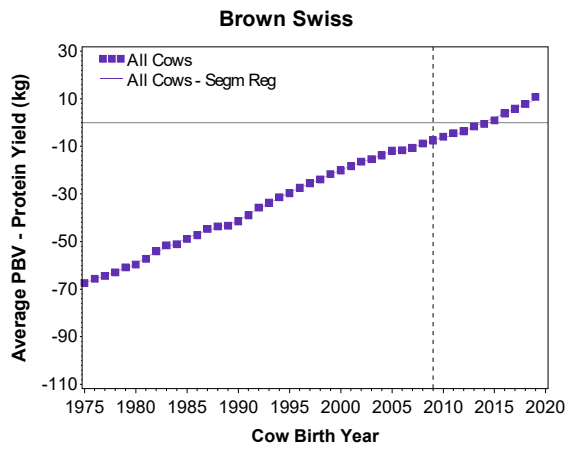
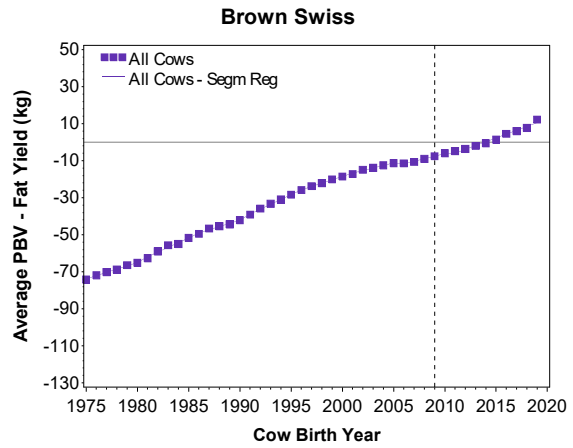
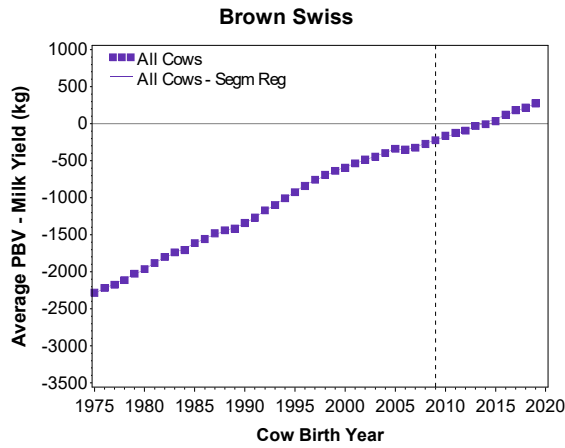


Figure S1. Estimates of genetic change per year from segmented regressions of \overline{PBV} on birth year for AY cows for seven traits: milk, fat, and protein yields; SCS; PL; DPR and LIV. The vertical dashed line indicates the year when genomic selection was implemented (i.e., 2013).



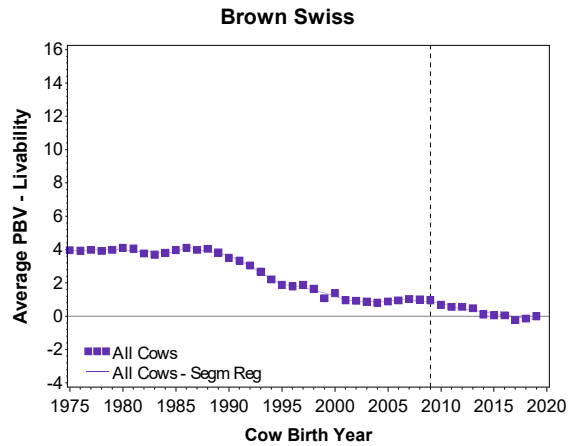
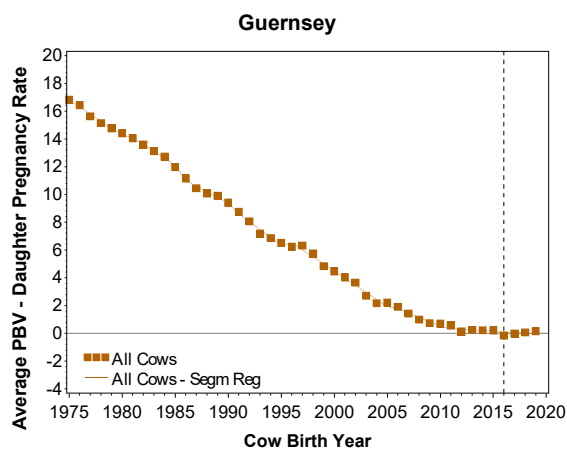
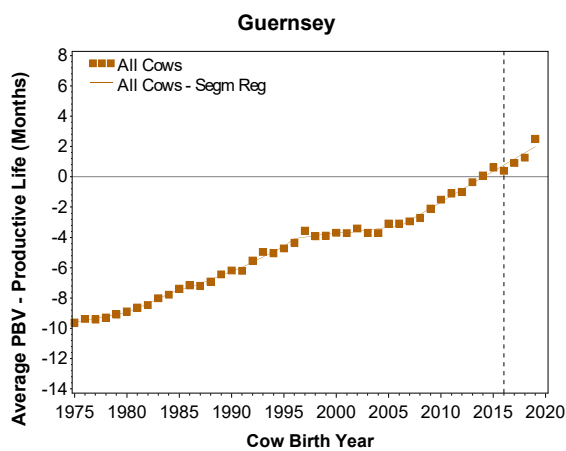
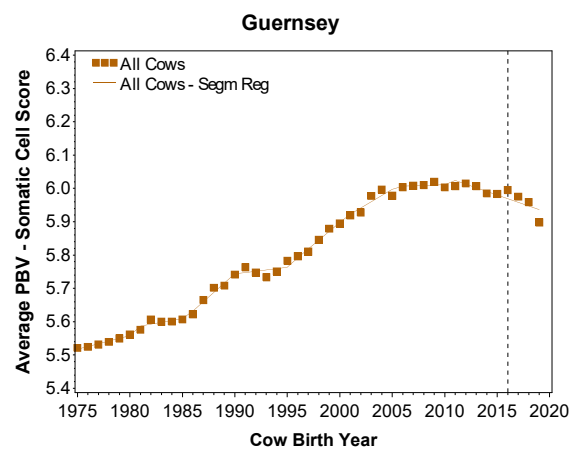
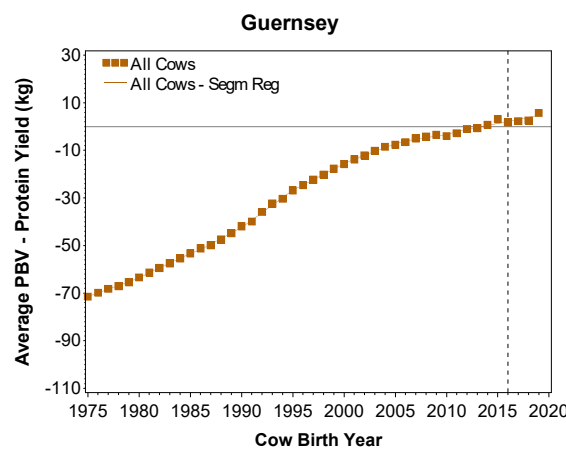
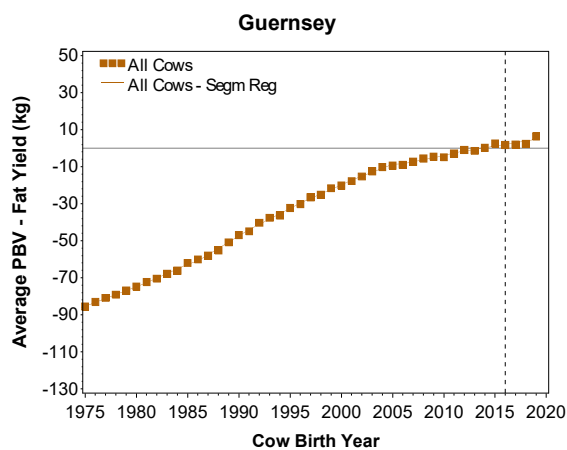
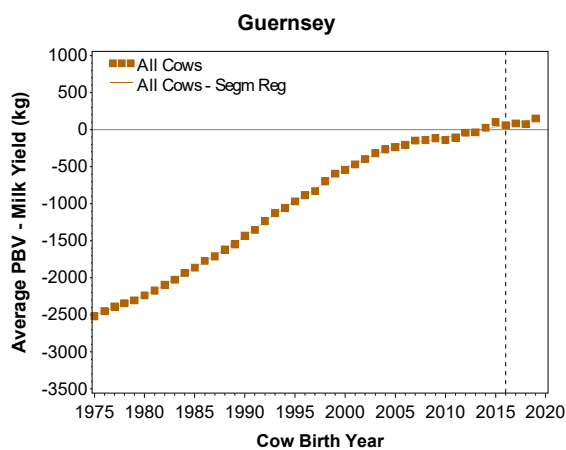


Figure S2. Estimates of genetic change per year from segmented regressions of \overline{PBV} on birth year for BS cows for seven traits: milk, fat, and protein yields; SCS; PL; DPR and LIV. The vertical dashed line indicates the year when genomic selection was implemented (i.e., 2009).



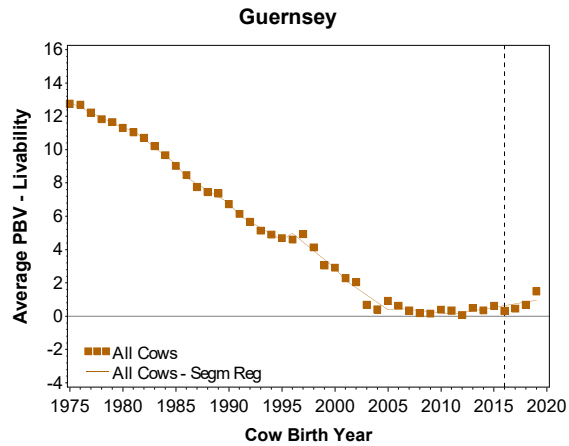
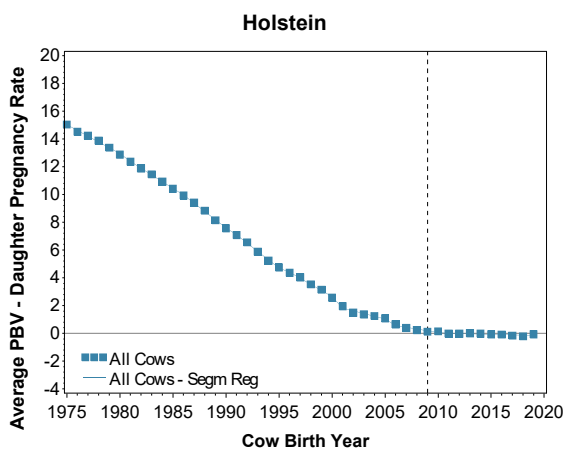
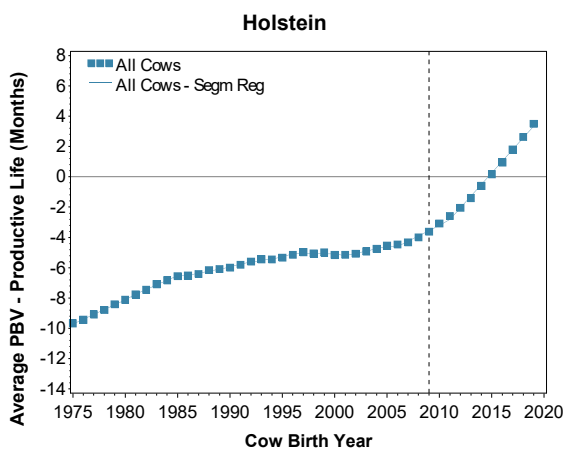
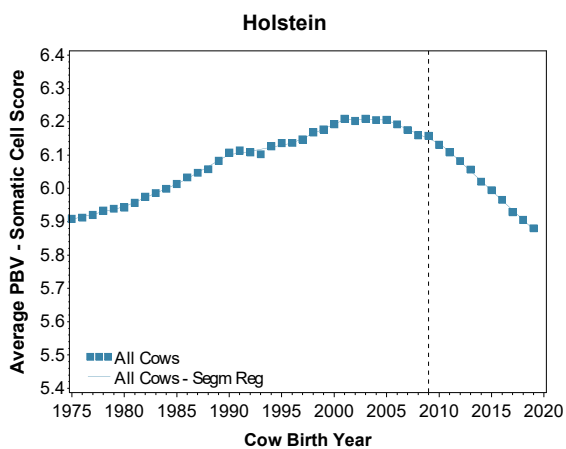
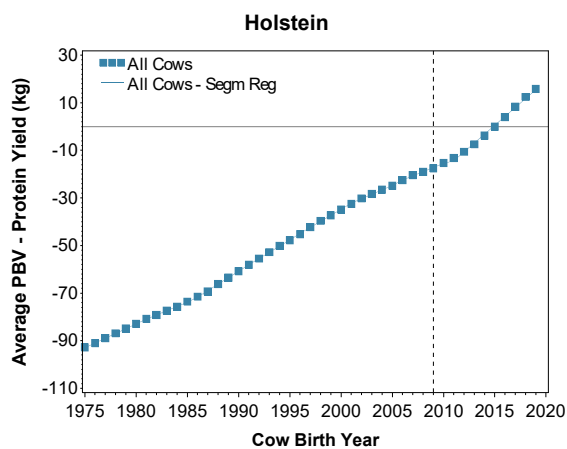
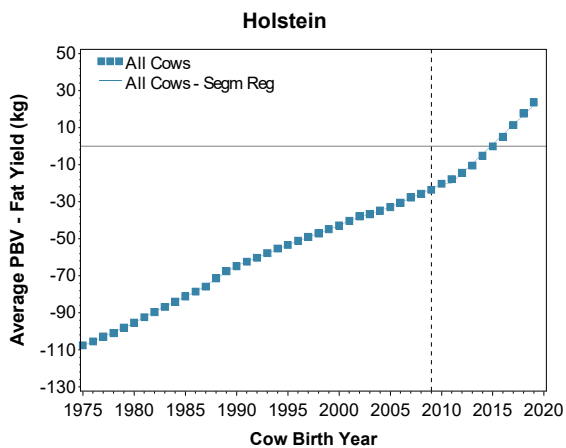
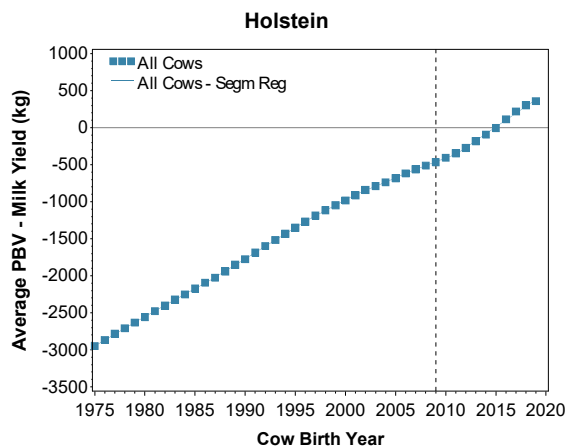


Figure S3. Estimates of genetic change per year from segmented regressions of \overline{PBV} on birth year for GU cows for seven traits: milk, fat, and protein yields; SCS; PL; DPR and LIV. The vertical dashed line indicates the year when genomic selection was implemented (i.e., 2016).



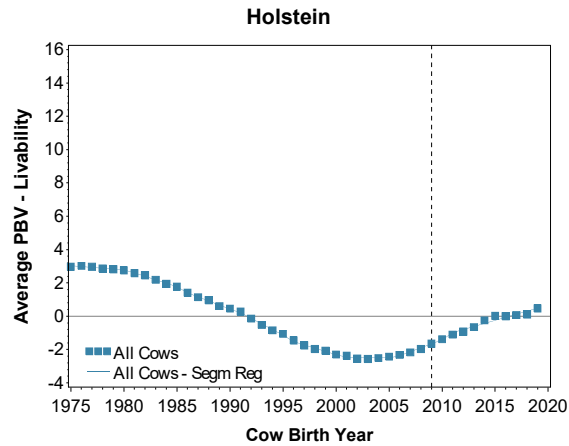
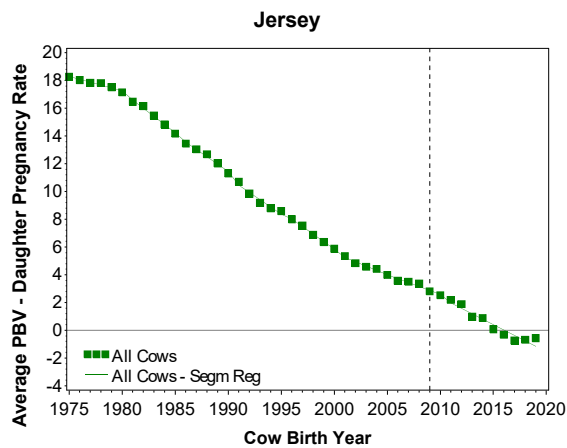
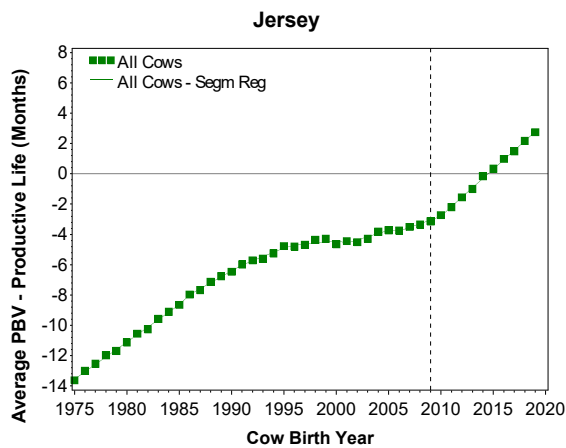
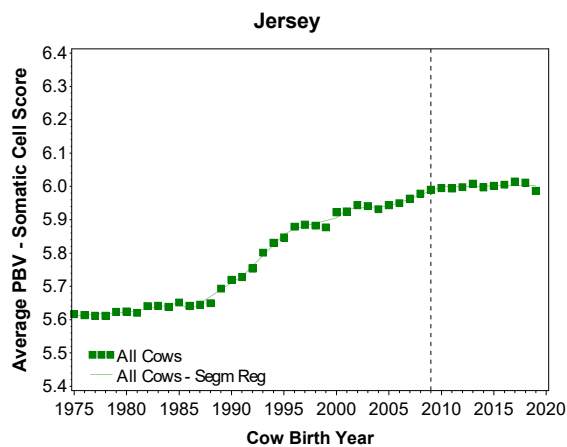
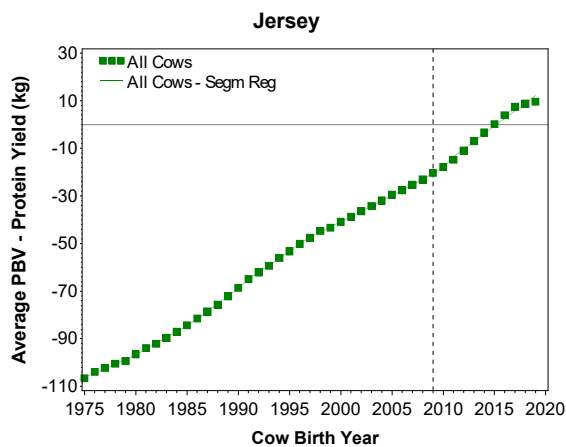
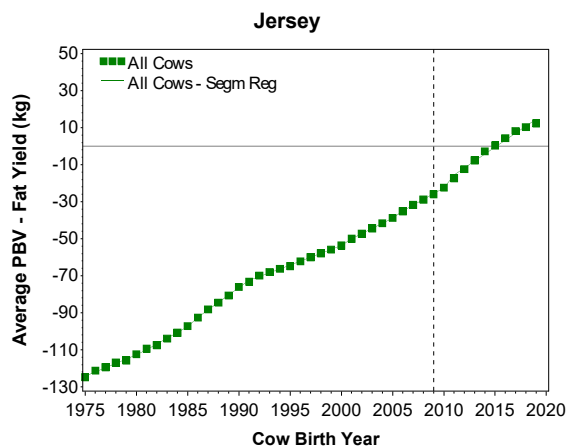
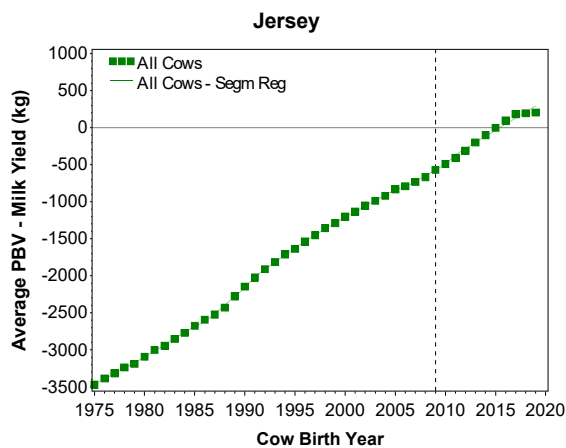
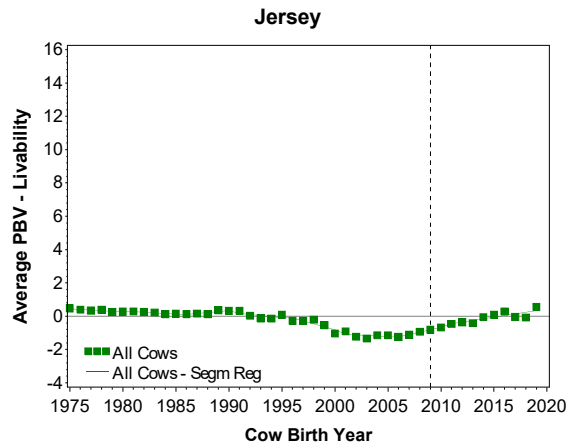


Figure S4. Estimates of genetic change per year from segmented regressions of \overline{PBV} on birth year for HO cows for seven traits: milk, fat, and protein yields; SCS; PL; DPR and LIV. The vertical dashed line indicates the year when genomic selection was implemented (i.e., 2009).





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Figure S5. Estimates of genetic change per year from segmented regressions of \overline{PBV} on birth year for JE cows for seven traits: milk, fat, and protein yields; SCS; PL; DPR and LIV. The vertical dashed line indicates the year when genomic selection was implemented (i.e., 2009).