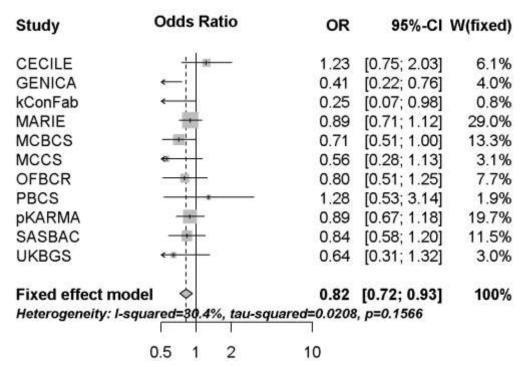
Supplementary Figure 6. Forest plot for SNPxHRT interactions regarding overall breast cancer risk by study for (A) rs9578047, (B) rs9579199, (C) rs7148646, (D) rs848694, (E) rs7192724, (F) rs17202296, and (G) rs4888190

A rs9578047



B rs9579199

Study	Odds Ratio	OR	95%-CI	W(fixed)
CECILE	; a	1.23	[0.75; 2.03]	6.1%
GENICA	<	0.41	[0.22; 0.76]	4.0%
kConFab	< i	0.25	[0.07; 0.98]	0.8%
MARIE		0.89	[0.71; 1.12]	29.0%
MCBCS		0.72	[0.51; 1.01]	13.3%
MCCS	a 🕂	0.56	[0.28; 1.13]	3.1%
OFBCR		0.80	[0.51; 1.25]	7.6%
PBCS		1.28	[0.53; 3.14]	1.9%
pKARMA	- 100	0.89	[0.67; 1.18]	19.7%
SASBAC	- (8)	0.84	[0.58; 1.20]	11.5%
UKBGS	<	0.64	[0.31; 1.32]	3.0%
Fixed effect m	nodel 🔶	0.82	[0.72; 0.93]	100%
Heterogeneity:	I-squared=30.3%, tau-squ	uared=0.0208	3, p=0.1573	
	0.5 1 2	10		

C rs7148646

Study	Odds Ratio	OR	95%-CI	W(fixed)
GENICA	~ + -	0.78	[0.45; 1.36]	6.7%
kConFab	<	0.22	[0.04; 1.27]	0.7%
MARIE		0.76	[0.61; 0.95]	39.3%
MCBCS		0.95	[0.66; 1.37]	15.5%
MCCS	«	0.66	[0.35; 1.24]	5.0%
OFBCR		0.84	[0.55; 1.27]	11.7%
PBCS	< + 	0.70	[0.28; 1.74]	2.4%
SASBAC	- **	1.03	[0.70; 1.52]	13.6%
UKBGS	<=	0.60	[0.32; 1.12]	5.1%
Fixed effect n	nodel 🔶	0.81	[0.70; 0.93]	100%
Heterogeneity:	I-squared=0%, tau-square	ed=0, p=0.63	04	
	0.5 1 2	10		

D rs848694

Study	Odds Ratio	OR	95%-CI	W(fixed)
GENICA	<u>← m¦</u>	0.77	[0.42; 1.40]	5.1%
kConFab	~ i	0.28	[0.05; 1.71]	0.6%
MARIE		0.77	[0.61; 0.99]	30.9%
MCBCS	- 181	0.86	[0.59; 1.27]	12.5%
MCCS	~ * :	0.72	[0.36; 1.42]	3.9%
OFBCR	<u>← mi</u>	0.76	[0.49; 1.18]	9.5%
PBCS	<	0.84	[0.33; 2.12]	2.1%
pKARMA		0.99	[0.73; 1.33]	20.5%
SASBAC		0.96	[0.64; 1.44]	11.1%
UKBGS	< <u>*</u>	0.90	[0.46; 1.77]	4.0%
Fixed effect n	nodel 🔶	0.84	[0.74; 0.96]	100%
Heterogeneity:	l-squared=0%, tau-squar	ed=0, p=0.91	42	
	0.5 1 2	10		

E rs7192724

Study	Odds Ratio	OR	95%-CI	W(fixed)
GENICA		1.28	[0.72; 2.28]	5.6%
kConFab	< i	- 1.23	[0.27; 5.54]	0.8%
MARIE	-	1.16	[0.91; 1.50]	29.5%
MCBCS		1.08	[0.75; 1.55]	14.1%
MCCS		1.11	[0.54; 2.30]	3.5%
OFBCR		1.45	[0.91; 2.31]	8.5%
PBCS	•	0.56	[0.20; 1.55]	1.8%
pKARMA		1.23	[0.91; 1.66]	20.6%
SASBAC		1.64	[1.11; 2.43]	12.2%
UKBGS	+ -	1.90	[0.91; 3.97]	3.4%
Fixed effect mo	del 🔶	1.25	[1.09; 1.43]	100%
Heterogeneity: I-s	quared=0%, tau-squar	ed=0, p=0.64	25	
	0.5 1 2	10		

F rs17202296

Study	Odds Ratio	OR	95%-CI	W(fixed)
GENICA		1.14	[0.66; 1.97]	5.9%
kConFab	< 	0.95	[0.25; 3.63]	1.0%
MARIE	-	1.01	[0.80; 1.29]	30.4%
MCBCS	- 100	1.15	[0.81; 1.64]	14.0%
MCCS		1.20	[0.59; 2.41]	3.5%
OFBCR	; .	1.50	[0.96; 2.36]	8.5%
PBCS	~ • 	0.77	[0.30; 2.02]	1.9%
pKARMA		1.00	[0.74; 1.35]	19.3%
SASBAC		1.43	[0.98; 2.09]	12.1%
UKBGS		1.42	[0.70; 2.89]	3.4%
Fixed effect m	nodel 🖕	1.13	[0.99; 1.29]	100%
Heterogeneity: I	-squared=0%, tau-squar	ed=0, p=0.79	01	
	0.5 1 2	10		

G rs4888190

Study	Odds Ratio	OR	95%-CI	W(fixed)
GENICA		1.24	[0.71; 2.14]	5.8%
kConFab	<u>← +</u>	0.96	[0.25; 3.64]	1.0%
MARIE		0.96	[0.75; 1.22]	30.4%
MCBCS		1.10	[0.77; 1.57]	14.0%
MCCS		1.45	[0.69; 3.05]	3.2%
OFBCR		1.36	[0.86; 2.16]	8.3%
PBCS	← + <u> </u>	0.83	[0.31; 2.17]	1.9%
pKARMA		0.95	[0.70; 1.28]	19.4%
SASBAC		1.35	[0.92; 1.97]	12.3%
UKBGS		1.37	[0.68; 2.76]	3.6%
Fixed effect mo	del 💠	1.09	[0.95; 1.24]	100%
Heterogeneity: I-s	quared=0%, tau-squar	ed=0, p=0.78	32	
		1		
	0.5 1 2	10		