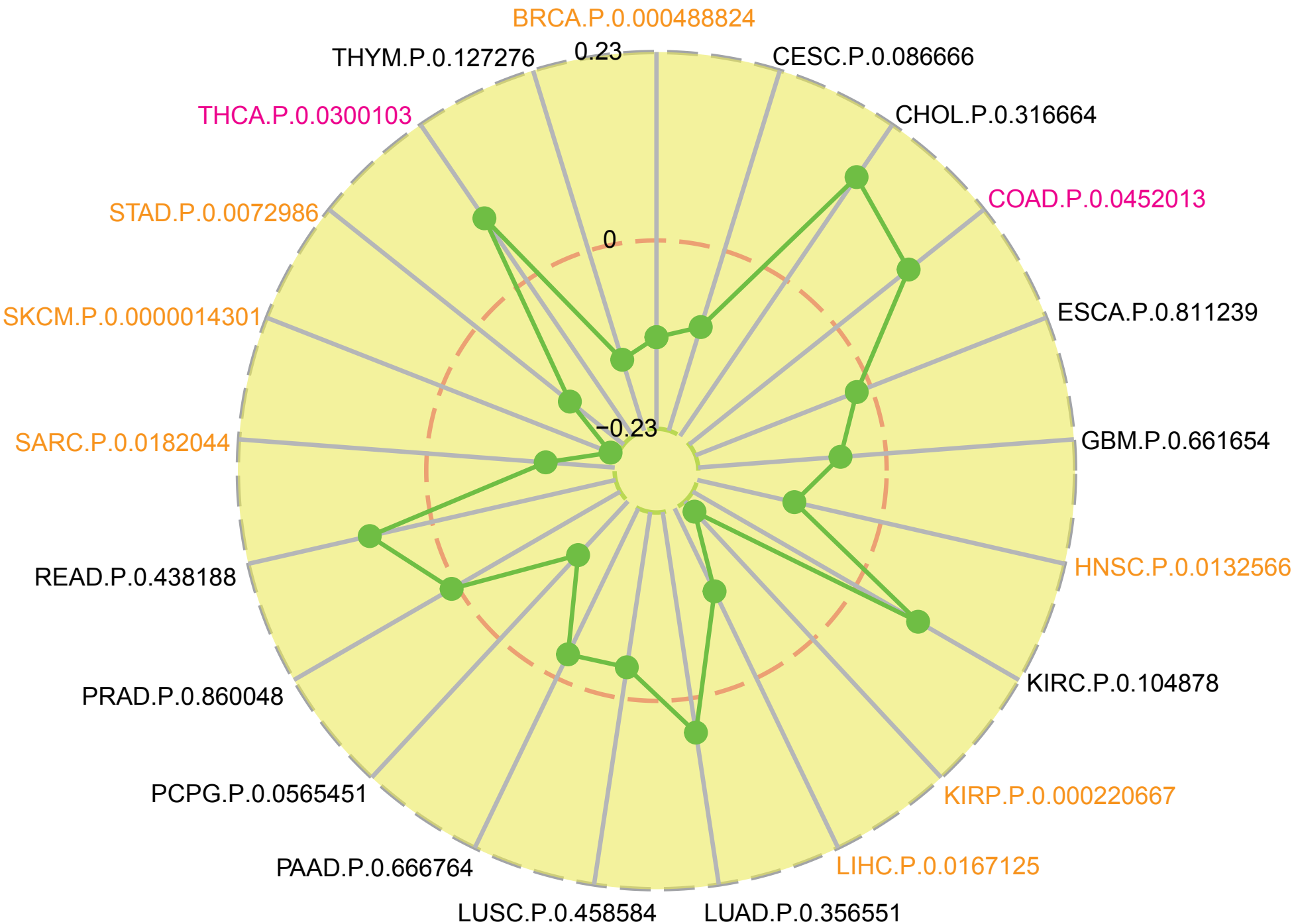Methylation−Expression correlation for KIF13A in pancancer