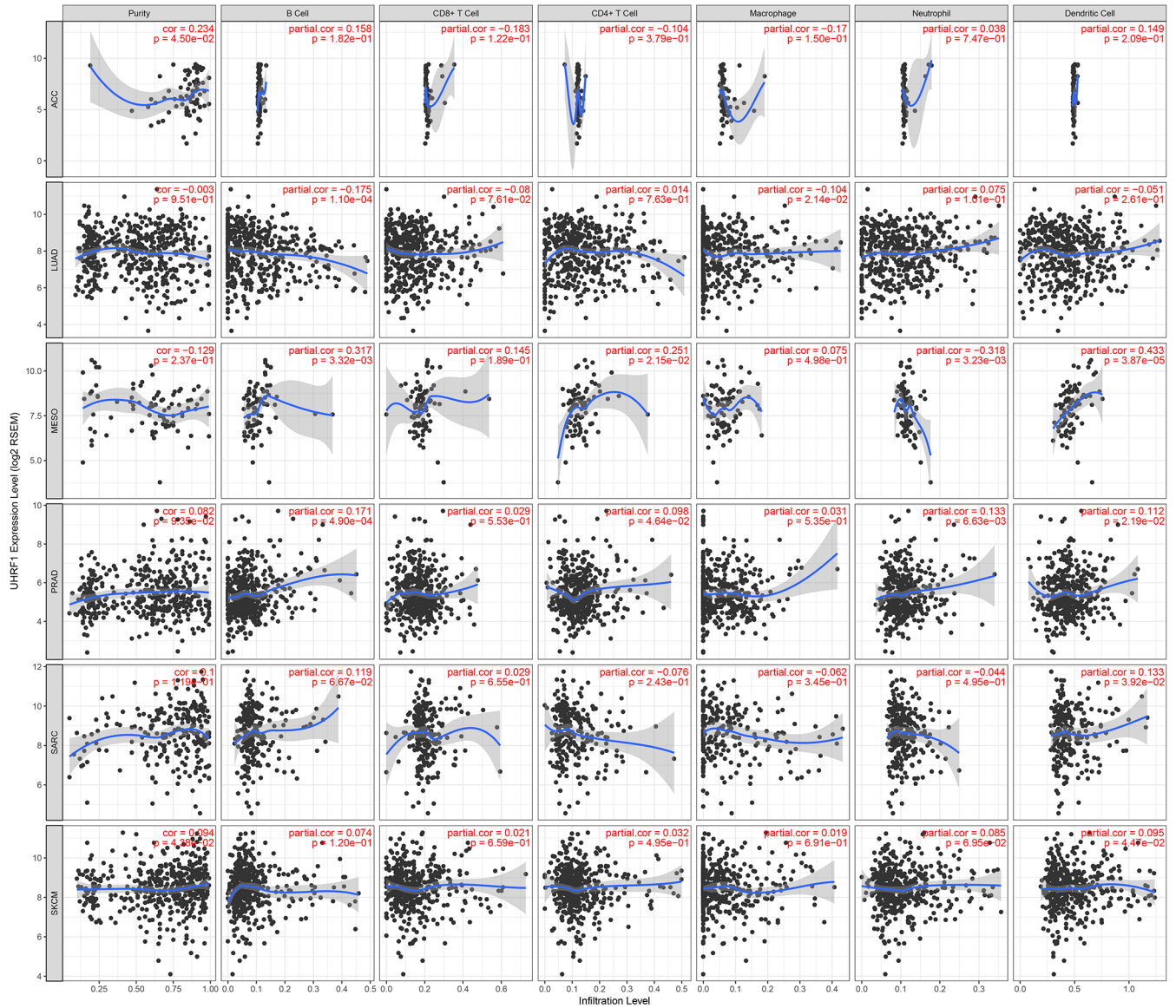


SUPPLEMENTARY FIGURES

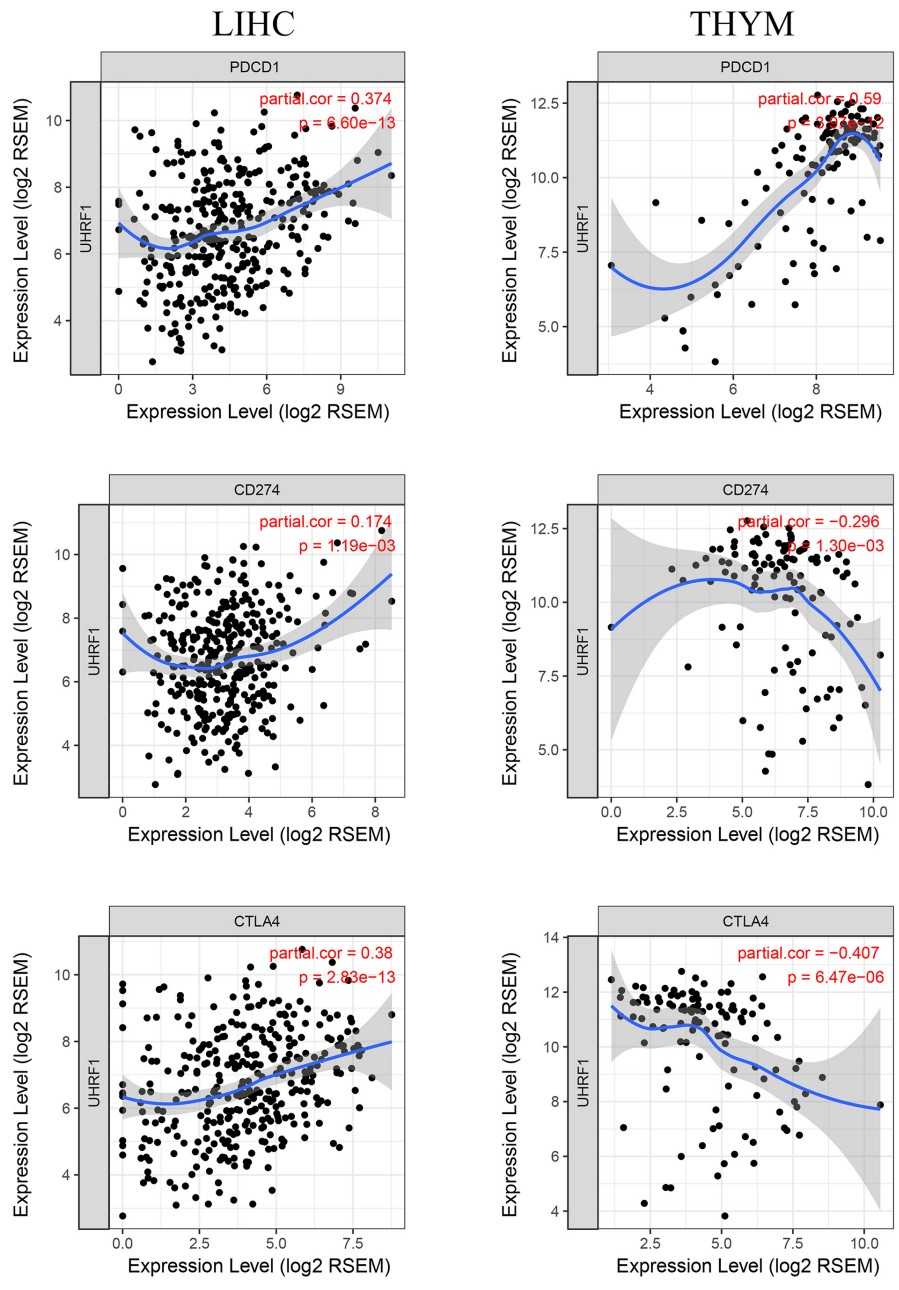
Range 1: 1197 to 2619 [GenBank](#) [Graphics](#) ▼ Next Match ▲ Previous Match

Score	Expect	Identities	Gaps	Strand
2584 bits(1399)	0.0	1415/1423(99%)	0/1423(0%)	Plus/Plus
Query 1	TGCGGGGGCCGGCAGGACCCCGACAAGCAGCTCATGTGCGATGAGTGCACATGGCCTTCCACATCTACTGCCTGGACCCGCCCTCAGC	90		
Sbjct 1197	TGCGGGGGCCGGCAGGACCCCGACAAGCAGCTCATGTGCGATGAGTGCACATGGCCTTCCACATCTACTGCCTGGACCCGCCCTCAGC	1286		
Query 91	AGTGTTCACAGCAGGACGAGTGGTACTGCCTGAGTGCAGGAATGATGCCAGCAGGTTGGTACTGGCGGGAGAGCGCTGAGAGAGAC	180		
Sbjct 1287	AGTGTTCACAGCAGGACGAGTGGTACTGCCTGAGTGCAGGAATGATGCCAGCAGGTTGGTACTGGCGGGAGAGCGCTGAGAGAGAC	1376		
Query 181	AAGAAGAAGGCAGAGATGGCCTCGGCCACATCGTCTCACAGCGGGACTGGGCAAGGGCATGGCCTGTGTGGCCGCACCAAGGAATGT	270		
Sbjct 1377	AAGAAGAAGGCAGAGATGGCCTCGGCCACATCGTCTCACAGCGGGACTGGGCAAGGGCATGGCCTGTGTGGCCGCACCAAGGAATGT	1466		
Query 271	ACCATCATCCCGTCCCAACCACTACGGACCCATCCCGGGGATCCCGTGGGCACCATGTGGCGGTTCCGAGTCCAGGTGAGCGAGTCCGGGT	360		
Sbjct 1467	ACCATCATCCCGTCCCAACCACTACGGACCCATCCCGGGGATCCCGTGGGCACCATGTGGCGGTTCCGAGTCCAGGTGAGCGAGTCCGGGT	1556		
Query 361	GTCATCGGGCCACGTTGGCTGGCATCCATGGCCGGAGCAACGAGCGGACTACTCCCTAGTCTGGCGGGGGCTACGAGGATGAGCTG	450		
Sbjct 1557	GTCATCGGGCCACGTTGGCTGGCATCCATGGCCGGAGCAACGAGCGGACTACTCCCTAGTCTGGCGGGGGCTACGAGGATGAGCTG	1646		
Query 451	GACCATGGGAATTTTTTTCACATACACGGGTAGTGGTGGTGCAGAGCTTTCCGGCAACAAGAGGACCCGGGAACAGTCTTGTGATCAGAAA	540		
Sbjct 1647	GACCATGGGAATTTTTTTCACATACACGGGTAGTGGTGGTGCAGAGCTTTCCGGCAACAAGAGGACCCGGGAACAGTCTTGTGATCAGAAA	1736		
Query 541	CTCACCAACACCAACAGGGCGCTGGCTCTCAACTGCTTTGCTCCCATCAATGACCAAGAAGGGCCGAGGCCAAGGACTGGCGTCCGGG	630		
Sbjct 1737	CTCACCAACACCAACAGGGCGCTGGCTCTCAACTGCTTTGCTCCCATCAATGACCAAGAAGGGCCGAGGCCAAGGACTGGCGTCCGGG	1826		
Query 631	AAGCCGGTCAGGGTGGTGCAGCAATGTCAAGGGTGGCAAGAATAGCAAGTACGCCCCGCTGAGGGCAACCGCTACGATGGCATCTACAAG	720		
Sbjct 1827	AAGCCGGTCAGGGTGGTGCAGCAATGTCAAGGGTGGCAAGAATAGCAAGTACGCCCCGCTGAGGGCAACCGCTACGATGGCATCTACAAG	1916		
Query 721	GTTGTGAAATACTGGCCCGAGAAGGGGAAGTCGGGTTTCTCGTGTGGCGCTACCTTCTGCGGAGGGACGATGATGAGCCCGGCCCTTGG	810		
Sbjct 1917	GTTGTGAAATACTGGCCCGAGAAGGGGAAGTCGGGTTTCTCGTGTGGCGCTACCTTCTGCGGAGGGACGATGATGAGCCCGGCCCTTGG	2006		
Query 811	ACGAAGGAGGGGAAGGACCGGATCAAGAAGCTGGGGTGCACATGCAGTATCCAGAAGGCTACCTGGAAGCCCTGGCCAACGgagagcga	900		
Sbjct 2007	ACGAAGGAGGGGAAGGACCGGATCAAGAAGCTGGGGTGCACATGCAGTATCCAGAAGGCTACCTGGAAGCCCTGGCCAACCGGAGAGCGGA	2096		
Query 901	gagaagagaaacagcaagagggaggaggaggagcagcaggaggggggCTTCGCGTCCCCAGGACGGGCAAGGGCAAGTGGAAAGCGGAAG	990		
Sbjct 2097	GAGAAGGAAACAGCAAGAGGGAGGAGGAGGAGCAGCAGGAGGGGGGCTTCGCGTCCCCAGGACGGGCAAGGGCAAGTGGAAAGCGGAAG	2186		
Query 991	TCGGCAGGAGGTGGCCCGAGCAGGGCCGGTCCCCCGCCGGACATCCAAGAAAACCAAGGTGGAGCCCTACAGTCTCACGGCCAGCAG	1080		
Sbjct 2187	TCGGCAGGAGGTGGCCCGAGCAGGGCCGGTCCCCCGCCGGACATCCAAGAAAACCAAGGTGGAGCCCTACAGTCTCACGGCCAGCAG	2276		
Query 1081	AGCAGCCTCATCAGAGAGGACAAGAGCAACGCCAAGCTGTGGAATGAGGTCTTGGCGTCACTCAAGGACCGGCCGGCAGCGCCAGCCCG	1170		
Sbjct 2277	AGCAGCCTCATCAGAGAGGACAAGAGCAACGCCAAGCTGTGGAATGAGGTCTTGGCGTCACTCAAGGACCGGCCGGCAGCGCCAGCCCG	2366		
Query 1171	TTCAGTTGTTCTGAGTAAAGTGGAGGAGACGTTCCAGTGTATCTGCTGTCAGGAGCTGGTGTTCGGGCCATCACGACCGTGTGCCAG	1260		
Sbjct 2367	TTCAGTTGTTCTGAGTAAAGTGGAGGAGACGTTCCAGTGTATCTGCTGTCAGGAGCTGGTGTTCGGGCCATCACGACCGTGTGCCAG	2456		
Query 1261	CACAACGTGTGCAAGGACTGCCTGGACAGATCCTTTCGGGCACAGGTGTTTTCAGTGCCTGCCCTGCCCTACGACCTGGGCCGAGCTAT	1350		
Sbjct 2457	CACAACGTGTGCAAGGACTGCCTGGACAGATCCTTTCGGGCACAGGTGTTTTCAGTGCCTGCCCTGCCCTACGACCTGGGCCGAGCTAT	2546		
Query 1351	GCCATGCAGGTGAACCAAGCCTCTGCAGACCGTCTCAACCAGCTCTTCCCGGGTACGGCAATGGCCGGTAT	1423		
Sbjct 2547	GCCATGCAGGTGAACCAAGCCTCTGCAGACCGTCTCAACCAGCTCTTCCCGGGTACGGCAATGGCCGGTAT	2619		

Supplementary Figure 1. Sequence similarity between RP11-424C20.2 and its parental gene UHRF1 (NM_001048201.2).



Supplementary Figure 2. Correlation of UHRF1 expression with immune infiltration in ACC, LUAD, MESO, PRAD, SARC and SKCM analyzed using the “Gene” module in TIMER.



Supplementary Figure 3. Correlation analysis between UHRF1 expression and PD-1, PD-L1 and CTLA-4.