S1 Figure

CLUSTAL Omega (version 1.2.4) multiple sequence alignment of the low-affinity and high affinity sites of human NRF2 (bold) which bind to the Keap 1 propeller protein and proposed similar sites in the *P. falciparum* WD40 repeat protein.

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NRF2        QQDMDLIDILWRQDIDLGVSR-EVFDFSQR----------------RKEYELEKQKKL-- 54
PfW7K5T2    DKGLNSFIILWKINETSFLDPIKKFYYNCEENEKKRTHIYNNKKKNQDEYDINQINDNTI 177
            ::.:: : ***: :    :.  : * :. .                 :.**:::: :.

NRF2        EK--ERQE---------QLQKEQ----------EKA-FFAQLQLDEETGE 82
PfW7K5T2    DNSSENGDAINNNVGYECIDISHDNRYICALTEEKIYIKSNTKYDDETGE 227
            ::  *. :          :: .:          **  : :: : *:****
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In NRF2 there are seven lysine **(K)** residues between the 2 sites, six of which are destined to be ubiquitinated in the propeller. In the *P. falciparum* WD40 repeat protein there are six Lys residues between the putative low and high affinity sites. A closely similar picture for putative low and high affinity sites in WD40 was confirmed in *P. reichenowi* and *P. gaboni*, although sufficient sequence was not available in the PlasmoDB database to confirm the similarity of the proposed high affinity binding site in other species.