# Validation and Fitness Testing of a Quantitative Immunoassay for HIF1 $\alpha$ in Biopsy Specimens



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## Abstract (#C3)

HIF1 $\alpha$  is an important marker of hypoxia in human tumors that is altered in a number of cancers, but a reproducible method to determine changes in HIF1 $\alpha$  protein expression in human tumor biopsy specimens has not been available. HIF1 $\alpha$  is being used as a pharmacodynamic marker in clinical trials in the NCI (CTEP# 8610, 9534). However, an important limitation for studying the response of HIF1α to cancer therapeutic agents is the lability of the protein, in the presence of oxygen, upon collection of the sample. We have devised a method of specimen collection, handling, and extraction that preserves and stabilizes HIF1 $\alpha$  levels in tumor biopsies. Employing this specimen-handling method allowed validation of a two-site immunoassay for HIF1 $\alpha$  quantitation in solid-tissue extracts, such as tumor biopsies. Intra-assay variability was less than 10%, and inter-assay variability was less than 20%. Accuracy, assessed by spike recovery, was 99 +/- 7%. HIF1 $\alpha$  readings declined linearly with decreasing sample load over a range of 1 to 10 µg protein per well. Fitness for purpose was demonstrated by quantifying a reduction in HIF1 $\alpha$  protein levels following topotecan treatment of a xenograft model. HIF1α was also demonstrated to be upregulated under low oxygen tension culture conditions in DU145 human prostate cancer cells. The HIF1 $\alpha$  immunoassay is currently being transferred to the NCI's National Clinical Target Validation Laboratory for use in support of NCIsponsored early clinical trials. This research has been funded with federal funds from the National Cancer Institute, National Institutes of Health, under Contract No. HHSN261200800001E.

### Materials and Methods

- The assay is a two-site chemiluminescence readout enzyme immunoassay. The assay uses a purified monoclonal antibody to HIF1 $\alpha$  (R&D Systems, 841689) as the capture reagent and a biotin-labeled goat anti-HIF1α polyclonal antibody (R&D Systems, 841690) as a probe, followed by streptavidin conjugated to horseradish peroxidase (HRP) (R&D Systems, 890803) as the reporter. Prior to the addition of tumor lysate, the extracts are assayed for total protein using the BCA assay (Pierce), and then diluted in Reagent Diluent (5% BSA in Wash Buffer – 1X PBS, pH 7.2–7.4 with 0.05% vol/vol Tween) to a concentration of 1  $\mu$ g/ $\mu$ L for the assay. 10  $\mu$ L of specimens diluted in Reagent Diluent are added to wells containing 90 µL of Reagent Diluent (10  $\mu$ g/well). All specimens are assayed in duplicate or triplicate. Capture antibody and specimen incubation are 16 + - 1 hr at 2-8°C.
- Standard is purified recombinant human HIF1α (amino acid residues 575-826; R&D Systems, 841691) diluted in Reagent Diluent. Assay dynamic range is approximately 7.8 to 1,000 pg/mL. The assay controls are produced from HIF1 $\alpha$ -transfected HEK293 (1  $\mu$ g/ $\mu$ L) (OriGene Technology, Inc., Cat # LY419880) and PC3 cell extract (1  $\mu g/\mu L$ ). The PC3 cells are harvested after washing in cold PBS containing protease inhibitor and lysed in cell extraction buffer (CEB), and processed as for cell extracts. The PC3 extracts are assayed for total protein using the BCA assay (Pierce) and then diluted to a concentration of  $1 \mu g/\mu L$  for the assay. High-control is a mixture of 0.5 μg of HEK293 and 9.5 μg of PC3 per well. Medium-control is a mixture of 0.125 μg of HEK293 and 9.875 µg of PC3 per well. Low-control is PC3 10 µg per well. Dilutions are made in Reagent Diluent to achieve reads in the desired range of the assay, and expected ranges of assay readouts for the controls are provided for each lot. High-controls will read 683–869 pg/ml HIF1 $\alpha$ , while mid-range controls will read in the range of 205–313 pg/ml and low controls will read between 41 and 69  $pg/ml HIF1\alpha$ .

#### **Degassing buffer:**

- 1. Cell extraction buffer was prepared on ice with protease inhibitor.
- 2. 400 ul of CEB with protease inhibitor was added to the microfuge tube with O-ring screw cap and kept on ice.
- 3. Nitrogen gas was gently bubbled through the buffer at the bottom of the microfuge tube using a pipette tip until the bubble replaced the air above the buffer. It takes about 5–10 sec to replace air in the tube with nitrogen gas.
- 4. The vial was sealed immediately by closing the cap tightly.
- 5. The buffer was shipped to NCTVL on ice for biopsy collection.
- 6. The buffer can be stored for a month at 2–8°C.

#### **Biopsy sample collection:**

- 1. Mix the biopsy collection tube containing degassed buffer by inverting several times.
- 2. Prepare 3–4 vials for collecting samples at a time.
- 3. Keep the vials upright in ice bath.
- 5. Place needle biopsy into the vial by placing the tip of the biopsy needle **into the buffer** in the cone at the bottom of the vial, then ejecting the tissue core.
- 6. Seal the vial immediately by closing the cap tightly.
- 7. Flash-freeze the vials for storage.

#### **Specimen Collection and Processing Are Critical to** Successful Development of HIF1α Immunoassay

- Prepare cell extraction buffer (CEB) on ice:
  - Protease inhibitor (Roche) 1:50

Components	MW	For 500 ml CEB
50 mM Tris	121.14	3028.5 mg
300 mM NaCl	58.44	8766 mg
10% Glycerol	92.09	50 ml
3 mM EDTA	0.5 M liquid	3 ml
1 mM MgCl <sub>2</sub>	95.22	47.5 mg
20 mM beta Glycerol	306.11	3061.2 mg
25 mM NaF	41.99	524.75 mg
1% Triton		5 ml
100 μM 2-hydroxyglutarate	192.08	9.604 mg

Add PMSF 1:1000 and Protease Inhibitor 1:50 to required volume.



Collection of cells in degassed buffer, plus processing of specimens in the hypoxia chamber yielded HIF1 $\alpha$ .



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- Specimen and control samples should be processed as follows:
- Make protease inhibitor cocktail by dissolving 1 tablet into 1 mL water
  - Fresh PMSF (100 mM stock solution, Sigma) 1:100



### HIF1α Expression in Immunoassay (PC3 Cells)

- Discrepancy between quantitative ELISA readout and Western blot: • Western says no HIF1 $\alpha$  in normoxic condition, and ELISA says it's there, but
- hypoxic is 3X higher. The Western says the differential is much greater than 3X. • Tissue processing a critical factor:
- Sonication is better than grinding with a homogenizer. • The data suggest that stabilization of HIF1 $\alpha$  during the assay is also critical.



### Assay Variability & Accuracy by Spike Recovery



#### Comparison of DU-145 Intratumoral HIF1α Levels in PMA-treated Mice

DU145 xenografts staged to 250 or >600 mg size. Treatment was phorbol 12-myristate 13-acetate [PMA] IP, QDx3. Specimen collection was by biopsy of live nude mice (N=6/treatment). Deep vs. shallow biopsies and degassed buffer followed by flash-frozen biopsy (Buffer) vs. flash-frozen biopsy alone (BTB) were compared.



There was no detectable effect of PMA injection on recovered HIF1 $\alpha$  levels. However, the degassed collection buffer in the collection vials yielded a significant increase of recovered HIF1 $\alpha$  levels compared to the BTB standard method (unpaired 2-tailed t-test = 0.004).

F1α	Spike Recover	У	
	EXPECTED		
	pg/mL	% RECOVERY	
	2001	95.2	
	501	88.9	
	126	112.6	
	2000	100	
	500	101	
	2006	103.8	
	506	105.9	
	131	96.2	
	125	92	
	2006	99.8	
	506	97.1	
		Mean 99	.30%
		SDEV 6	60%
	Mean (pg/mL HIF1	α) SDEV	CV%
ility			
	27.9	1.9	6.5
lity			
	29.3	2.0	6.7

#### Bead Homogenizer vs. Sonication (10 µg load)

#### Note that topotecan treatment significantly reduced HIF1 $\alpha$ expression vs. vehicle



\* Less than 10  $\mu$ g of protein was loaded due to the diluted sample concentration to <1  $\mu$ g/mL.

#### CEB vs. CEB + 2-HG (100 $\mu$ M) vs. $CEB + \alpha - KG (100 \ \mu M) (10 \ \mu g \ load)$



#### CEB vs. CEB + 2-HG (100 $\mu$ M) vs. CEB + $\alpha$ -KG (100 $\mu$ M) (10 $\mu$ g load) **Second Experiment**



These studies were all performed with archived specimens collected by the BTB standard method





**Experiment Design** 

- Animal: A375 (human melanoma) nu/nu mouse xenografts • Treatment:
- Vehicle (G1) Indenoisoquinoline (NSC 743400) 25 mg/kg QD x1 IV (G2) NSC 743400 25 mg/kg QD x5 IV (G3) NSC 743400 12.5 mg/kg QD x5 IV (G4) NSC 743400 2.5 mg/kg QD x5 IV (G5) Topotecan 4 mg/kg QDx5 IP (G6)
- Tumor collection: 1 hr post-dose 5
- Entire tumor was collected, divided into 4 pieces, and flash-frozen.
- Each tumor piece was divided into 4 pieces and was lysed in degassed buffer.
- Different methods of homogenization were compared between sonication and bead homogenization with 3 different beads (ceramic 1.4 mm, ceramic 2.8 mm, metal 2.8 mm).
- Different lysis buffers were compared (CEB vs. CEB + 2-HG 100 μM vs. CEG +  $\alpha$ -Ketoglutarate 100  $\mu$ M).
- 10 µg of protein was loaded in each well.





• Sampling time: 1 hr post-dose 5

## Conclusions

- HIF1α immunoassay was developed and shown to be useful in recovering HIF1 $\alpha$  levels in xenograft models.
- Alternative methods of homogenization were evaluated: Sonication and homogenization with ceramic beads were superior to homogenization with metallic beads or tissue grinding.
- Degassed collection buffer yielded a significant increase of recovered HIF1 $\alpha$  levels compared to the standard method of fresh-freeze only.
- Alternative collection buffers show a trend for improved yield with the addition of 2-hydroxyglutarate.
- Topotecan and indenoisoquinoline NSC 743400 significantly reduced the amount of HIF1 $\alpha$  in A375 xenografts.

All animals used in this research project were cared for and used humanely according to the following policies: the U.S. Public Health Service Policy on Humane Care and Use of Animals (2000); the Guide for the Care and Use of Laboratory Animals (1996); and the U.S. Government Principles for Utilization and Care of Vertebrate Animals Used in Testing, Research, and Training (1985). All NCI-Frederick animal facilities and the animal program are accredited by the Association for Assessment and Accreditation of Laboratory Animal Care Internationa





