

M	model	dur.	% atrophy or force decline*	lipid per. & protein ox.	ROS produc.	antioxid. defense systems	protein synt. & degr.	A	reference
<b>animals</b>									
dia	MV, rat	18 h	15-30	↑LP, ↑PO			↑PD, ↑calp, ↑UP		(Shanely <i>et al.</i> , 2002)
dia	MV, rat	6, 12, 18 h					↓PS		(Shanely <i>et al.</i> , 2004)
dia	MV, rat	12 h			↑DCFH	↓GSH, ↓GPX, ↑HO-1, ↓SOD			(Falk <i>et al.</i> , 2006)
dia	MV, rat	12 h	~15*(force)			↓GSH ↓thiols	↑PD		(Betters <i>et al.</i> , 2004)
			=			↓GSH ↓thiols	=PD	tro	
dia	MV, rat	18 h	~40	↑PO		↓GSH	↑UP (MuRF-1 & atr-1)		(McClung <i>et al.</i> , 2007)
			~10	=PO		↓GSH	↑UP (MuRF-1 & atr-1)	tro	
dia	Mv, rat	12 h	~20-30 ~20*(force)	↑PO, ↑LP			↑calp, ↑casp		(Whidden <i>et al.</i> , 2010)
			=	no PO, no LP			=calp, =casp	tro	
sol	imm, rat	4, 8, 12 d	20-50 (rate)	↑LP (60%)		↑GSSX/GSH, ↑ Fe			(Kondo <i>et al.</i> , 1992)
		12 d	~35 (rate)	↑LP (20%)				vit E	

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sol	imm, rat	4, 8, 12 d	19-51			↑SOD, ↓MnSOD, =Se-GPX, =cat, ↑GST, ↑GSHRx, ↑XOD			(Kondo <i>et al.</i> , 1993)
sol	HU, rat	28 d	45		↑DCFH, ↑tot hydr	↑SOD, ↓MnSOD, ↓cat, ↓GPX, ↓ASC			(Lawler <i>et al.</i> , 2003)
sol	HU, rat	28 d	55		↑ tot hydr	↓GPX, ↓Hsp70, ↓Hsp25			(Lawler <i>et al.</i> , 2006)
sol	imm, rat	8 d	35			↓GSH (small)			(Appell <i>et al.</i> , 1997)
			12			=GSH		vit E	
sol & gas	HU	14 d	44 (sol) 30 (gas) 28* (force)	↑PO, ↑LP					(Koesterer <i>et al.</i> , 2002)
			~44 (sol) ~30 (gas) ~28* (force)	↑PO, ↑LP				vit E	
sol	HU, mice	12 d	44-38 64*(force)		↑ox act				(Matuszczak <i>et al.</i> , 2004)
			44-38 *force recovers					AI	
sol	HU, mice	12 d	~40		↑DCFH		=ser prot		(Arbogast <i>et al.</i> , 2007)
			~15		=DCFH		↓ser prot	BBIC	

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gas	HU, rat	5-10 d	~20	↑LP		↓GSH, ↑GSSG, ↓GSH/GSSG	↑UP		(Ikemoto <i>et al.</i> , 2002a)
			partial restoration	=LP		partial restoration of GSH, GSSH, GSH/GSSG,	=UP	cyst	
gas	HU, rat	2-21 d	~10	↑LP		↓GSH, ↑GSSG	↑UP		(Ikemoto <i>et al.</i> , 2002b)
			~10	↑LP		↓GSH, ↑GSSG	↑UP	vit E	
sol	HU, rat	14 d	49	↑LP		↓GSH/GSSG, ↑SOD, ↑GPX, ↑cat	↑casp-3-9, ↑atr-1, ↑MuRF-1		(Servais <i>et al.</i> , 2007)
			32	↓LP		↓GSH/GSSG, ↑SOD, ↑GPX, ↑cat	↓casp-3-9, ↓atr-1, ↓MuRF-1	vit E	
sol	HU, mice	14 d	22	↑LP, ↑PO		↑SOD, ↓CAH-III, ↓PRX-6, ↓Hsp <sub>s</sub>			(Brocca <i>et al.</i> , 2010)
			22	=LP, =PO		↑SOD, ↑CAH-III, ↑PRX-6, ↑Hsp <sub>s</sub>		tro	
gas	HU, rat	14 d	12	↑LP	=H <sub>2</sub> O <sub>2</sub> , ↑nitr	=SOD, =MnSod, =cat			(Siu <i>et al.</i> , 2008)
gas	HU, mice	14 d	10	=LP, =PO		↑SOD, ↑MnSOD, ↑PRX-6, ↑CAH-III			(Brocca <i>et al.</i> , 2010)
			10	=LP, ↓PO		↑*CAH-III, ↑*PRX-6, ↑*Hsp60		tro	

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<b>humans</b>									
dia	MV	1-69 h	53-57			↓GSH, ↑casp	↑UP (atr-1, MuRF-1)		(Levine <i>et al.</i> , 2008)
VL	BR	20 d	3-5				↑UP, ↑MuRF-1, ↑atr-1,		(Ogawa <i>et al.</i> , 2006)
VL	BR	7, 35 d	0-18	=PO (8d), ↑PO (35d)		↑HO-1 (8d), =HO-1 (35d), ↑Grp75 (8d), =Grp75 (35d)			(Dalla Libera <i>et al.</i> , 2009)
VL	ULLS	10, 23 d	5-10				↓PS, =mTor, ↓MuRF-1, ↓atr-1		(de Boer <i>et al.</i> , 2007)
VL	imm	2, 14 d	5-6	=LP, =PO		↑UPc (2d), =UPc (14d) =casp (2-14d)			(Glover <i>et al.</i> , 2010)

Table 1. Summary of the key literature cited related to muscle atrophy and oxidative stress in order of appearance. Only major findings per reference are reported. Antioxidant treatments have gray background. All symbols refer to comparison with controls, i.e. between disuse atrophy and controls or between disuse atrophy following treatment and controls.

Abbreviations:

↑= higher than control	HO-1=heme oxygenase-1, heat shock protein-70 or mortalin
↓= lower than control	HU=hindlimb unloading
= unchanged with respect to control	imm= immobilization
A= antioxidant treatment	lipid per. =lipid peroxidation
AI = allopurinol	LP= lipid peroxidation
ASC= antioxidant scavenging capacity	M= muscle
atr-1= atrogin-1	MV= mechanical ventilation
BBIC= Bowman-Birk inhibitor	nitr= nitrothirosine content
BR= bed rest	ox act= oxidant activity
CAH=carbonic anhydrase	PD= protein degradation
calp= calpain	PO= protein oxidation
casp= caspase	protein ox.= protein oxidation
cat= catalase	protein synth & degr = protein synthesis and degradation
cyst= cysteine	PRX=peroxiredoxin
DCFH = dichlorofluorescin diacetate oxidation (index of ROS content)	PS= protein synthesis
dia= diaphragm	ROS produc.= ROS production
dur= duration	ser prot= serine proteases
gas= gastrocnemius	SOD= copper-zinc superoxide dismutase
GPX=glutathione peroxidase	sol= soleus
Grp-75= glucose-regulated protein-75 or mitochondrial	tot hydr= total hydroperoxides
GSH= glutathione	tro= trolox
GSHRx= glutathione reductase	UP= ubiquitin proteasome activity
GST= glutathione S-transferase	UPc= ubiquitin proteasome conjugates
	VL= vastus lateralis
	XOD= xanthine oxidase