The clinical problem: Implants for Joint replacements generate wear particles of different size distributions

Cytokines of macrophages after phagocytosis of nano- or micro-sized corundum, graphite and chromium oxide

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Cemented metal / metal total hip prosthesis after 25 years.

superficial necrosis of the neo-synovia with macrophages full of small particles. H&E.





Hydroxyapatite coated CoCrMo-alloy stem,  $Al_2O_3$  ceramic head, one year after implantation in the left femur, Male, 125 kg BW, E 13421-92



Ieft humero-acromial joint replacement 2 years, change of the prosthesis 3 months ago. Female 80 years. E 5148-80, polarized Nicols prisms.



# IH PGM1 macrophages, E 4444-91

	Fe	Cr	Ni	Co	
n	102	70	31	50	
max	5629	2646	739	2746	
min	101	0.7	3	4	
mean	936	197	64	239	

Content of Fe, Cr, Ni and Co in mg/kg or ppm in tissue adjacent to failed, surgically removed hip and knee prostheses, 79 cases. n = number of positive determinations



# Plasmapore coated Aesculap BiContact <sup>®</sup> stem

# Femur



### Background

Aseptic loosening of endoprostheses is mainly correlated to generation of particles

Particles are released from gliding surfaces of metal, metal alloy, UHMWPE, Alumina  $Al_2O_3$ , bone cement and components, e.g. zirconium

The size of particles varies from nm to µm

Particles are found in the interface, around the interface, adjacent tissues, lymph nodes, lung, spleen, liver and other organs

Particles are phagocytosed by macrophages and antigen presenting cells

Stimulation of cytokines in murine macrophages

# by micro- and nano-sized particles

**Macrophages** 

produce cytokines e.g. IL-1ß, IL-6, IL-10, IL-12, G-CSF, GM-CSF, TNF- $\alpha$ , KC, nitrogen oxide (NOx).

influence T-cell and B-cell response i.e. the immune response

sensitive against Lipopolysaccharide (LPS) derived from the external part of the wall of Gram- negative bacteriae

Lipid A anchored in the bacterial wall

## **Objectives**

Aluminium oxide polycrystalline ceramics as gliding components for hip and knee endoprostheses generate the smallest number of particles of nm and  $\mu$ m size.

From new very thin coatings of diamond-like carbon to metallic modular endoprostheses nm size carbon particles could be released.

Co-Cr-Mo alloys are used and chromium containing particles could be released.

**Question:** 

Do particles of nm or µm size generate different cytokines and NOx ?

# **Materials and Methods**

## **Characterization of materials**

(Wear 264 (2008 ) 505-517 JBMR-A Published Online: Apr 22 2008 ISSN: 1552-4965)

## Murine macrophages (RAW264.7 cells) ASTM standard F 1903

- Macrophages morphology, vitality, SEM, TEM
- Cytokines by Multiplex- ELISA
- NOx by Griess reagent
- Statistics ANOVA Man-Whitney test
- Gene expression profiles (mRNA) by TaqMan-PCR



µm MX51





#### chromium (III) oxide



particles	density [g/cm³]	Vol- conc.	40 ml sus- pension	D <sub>50</sub> [µm]	Zetapotent	ial [mV]	[mV] / pH			
		[Vol%]	(0,8 ml solid)		Cell culture medium	9	Dist. water			
					Zetapot.	рН	Zetapot.	рН		
corundum MR 52 (µm)	3,9442	2	3,19 g	1,7	+7,09	8,68	-14,16	8,68		
corundum LA (nm)	3,6768	2	2,94 g	0,04	+3,17	8,48	+7,56	7,79		
Cr <sub>2</sub> O <sub>3</sub> (nm)	5,0587	1	2,025	0,1	+0,66	8,0	-1,21	4,46		
graphite MX 51 (µm)	2,2735	2	1,82 g	6,3	+25,29	8,25	-10,4	10,33		
graphite (nm) Printex 90C	2,0380	1	0,815 g	0,1	-9,49	8,0	-3,25	8,6		

# Cytokine determination by Multiplex-ELISA (Bio-Plex, BioRad)



1. incubate antibody-coupled beads with sample or standard

2. wash, then incubate with biotinylated detection antibody

3. wash, then incubate with streptavidin-PE; read on the Bio-Plex suspension array system

IL-1ß, IL-6, IL-10, IL-12 (p40), IL-12 (p70), TNF-a, KC, G-CSF, GM-CSF

# **Results** cells RAW264.7, corundum









µm corundum 48 h

+ control LPS 48 h

nm corundum 48 h

# **SEM (RAW264.7)**







## -C, 48h, 2000x La c1, 48h, 2000x Mr c1 48h, 2000x







+C, 48h, 2000x La c2, 48h, 2000x Mr c2, 48h, 2000x

KT 001675 12E-10 M7000 <-->1.1vm RAW LAC1/48H US0-0000 I12E-10





nm particles КТ **001679 0**50607 М7000 <-->1.1ум RAW LAC1/96H U50-**0000** I12E-10



КТ 001674 12Е-10 М20000 <--> .4ум RAW LAC1/48H U80-0000 I13Е-10 KT 001678 050607 M7000 <-->1.12m RAW LAC1/96H U50-0000 I12E-10

attachment internalization phagocytotic vacuoles secondary lysosomes vacuoles as sign of stress

950607

LA/C148H

RAW

M7000

<-->1.1Pm

U50-0000 I12E-10

кт (	901677	12E-10	M12000	<	->	.6Pm
RAW	MRC1/4	вн	U80-000	00	112	E-10



КТ 001689 050607 M20000 <--> .4ум RAW MR/C148H U50-0000 I55E-11



KT 001685 050607 M12000 <--> .6Pm RAW MRC1/96H U50-0000 I12E-10



КТ 001671 12E-10 M3000 <-->2.5ум RAW+K48M U80-0000 I12E-10





+ and -

controls





## Gene profiling of 48 murine mRNAs

### by TaqMan-Low-Density-Array

Microbe-controlled

Inhibitory Signals

TO HUR DOS RECORDS

**Bio**Gipss

Migration/Motility

Antigen Presentation

Lections

Phagocyte Maturation

Actin

Inhibitory Signals

pathway



G-CSF
GM-CSF
IL-10
ll-1b
II-6
IL-12(p40)
MCP-1
Prostaglandin E Synthasse 2
TNF alpha
MIP-1a
IL15
M-CSF
IL1-Rezpetor
IL-1-receptor associated kinase1
TNF Receptor 1A (receptor 1)
TNF Receptor 1B (Receptor 2)
TRAF6
Ubiquitin
Rho GTPase (activating prot. 1)
Rho GTPase (activating prot. 4)
Rho GTPase (activating prot. 26)
Rho GTPase (activating prot. 29)
Cathepsin H
TLR4
NF kappa B1

(modified from D.M.Underhill and A.Ozinsky, Annu. Rev. Immunol. 2002. 20:825-852)

# Differentially regulated murine mRNAs after incubation of macrophages with different materials (24-72h incubation)



Quantification of differential gene expression of 48 murine mRNAs by TaqMan-LDA

			Receptor and	Nr. of tested materials
Nr.	Gene Symbol	Gene name	signalling interactions	influencing gene expression
1	Csf1	M-CSF	Cytokine	11
2	Nos2	iNOS	Oxidative Stress	10
3	Arhgap1	Rho GTPase (ad	IL-1 pathway	10
4	Csf3	G-CSF	Cytokine	10
5	Cr2	CD35 (CR2)	Complement	10
6	Dnm1	Dynamin 1	Complement	10
7	Mrc1	CD206	Phagocytose	9
8	ll12b	IL-12(p40)	Cytokine	9
9	Arhgap4	Rho GTPase (ad	IL-1 pathway	8
10	Arhgap26	Rho GTPase (ad	IL-1 pathway	7
11	li10	IL-10	Cytokine	7
12	ll1r1	IL1-Receptor	IL-1 pathway	6
13	Uchl1	Ubiquitin	IL-1 pathway	6
14	Ccl3	MIP-1a	Cytokine	6
15	ll1b	ll-1b	Cytokine	6
16	Fcgr3a	CD16(FCgRIII)	Phagocytose	5
17	Msr1	SR-A (macropha	Phagocytose	5
18	Ctsh	Cathepsin H	IL-1 pathway	5
19	Tnfrsf1b	TNF Receptor 1	IL-1 pathway	5
20	Traf6	TRAF6	IL-1 pathway	5
21	Cd68	CD68	Fc-Receptors	5
22	II15	IL15	Cytokine	5
23	Dnm2	Dynamin 2	Complement	5
24	Bax	Bax	Apoptose	5
25	Casp9	Caspase9	Apoptose	5

#### Levels of different protein concentration and mRNA level of selected cytokines

– μm- and nm-sized corundum after 72h incubation

TNF-α

G-CSF



### Levels of different protein concentration and mRNA level of selected cytokines

– µm- and nm-sized graphite after 72h incubation

										INF	-α	(	G-C	SF							
De	elta Ct (PC	R cycle	es) vs	. Hou	sekee	eping g	ene C	DK2		$\frown$			$\frown$								
			Csf2	ll10	Dnm1	Arhgap4	Ccl2	Arhgap26	Cc/3	Tnf	Uchl1	II12 <mark>1</mark> 5	Csf3	Mrc1	Csf1	Cr2	ll1r1	Mst1r	Arhgap1	ll1b	Nos2
24h	NC		-4.95	-0.40	-3.51	-0.17	0.79	-1.66	-2 <mark>/</mark> 26	0.12	3 44	-0.78	3.02	-9 <mark>0</mark> 7	-0.85	5.25	4.96	7.77	-0.42	-1.16	-2.49
24h	Graphite	MX51- µn	-4.11	1.06	0.88	-0.22	3.11	-1.91	1 <mark>.</mark> 89	1.29	5.00	2. <mark>∂</mark> 5	6.89	-5. <mark>7</mark> 6	2.34	6.02	2.12	3.02	3.02	-8.54	1.65
24h		P900- nm	-5.46	1.85	-0.12	-0.06	2.89	-0.36	1.85	1.05	4. <mark>3</mark> 5	1 <mark>4</mark> 1	5.79	-8.0 <mark>9</mark>	-0.52	8.93	3.91	2.55	0.75	-0.45	-0.56
72h	NC		-0.45	5.38	4.15	3.41	-1.49	8.80	(.91	-1.78	3. <mark>3</mark> 2	18 <mark>.19</mark>	12.77	10.39	11.30	19.06	11.02	11.77	17.45	6.44	6.92
72h	Graphite	MX51 - μ	-2.38	-0.12	0.92	-0.93	-0.07	0.34	( <mark>.44</mark>	-0.34	0. <mark>3</mark> 5	-3 <mark>5</mark> 2	-2.95	3.6 <mark>8</mark>	1.60	1.56	0.59	-3.69	-3.62	0.68	0.19
72h		P900 - nn	2.40	-0.24	1.49	-0.59	1.14	0.80	2 28	0.44	0 00	0.52	-7.92	2.78	2.78	-3.78	1.59	2.89	-1.82	-0.38	-0.69

## **TNF-α:** protein – 10fold

## G-CSF: protein – 2fold

### mRNA – 2fold





### mRNA – 32fold



#### Levels of different protein concentration and mRNA level of selected cytokines

- different concentrations of nm-sized chromium (III) oxide after 72h incubation



■ neg K ng/1E6 Zellen ■ pos K ng/1E6 Zellen ■ Cr2O3 ng/1E6 Zellen

■ neg K ng/1E6 Zellen ■ pos K ng/1E6 Zellen ■ Cr2O3 ng/1E6 Zellen

## Conclusions

Particles of different size distributions play a major role for failure of implant components.

In experiments with a macrophage cell line (RAW264.7) stimulated by model particles of corundum, graphite and  $Cr_2O_3$  with nm and  $\mu$ m size and identical mass, different genes were up- or downregulated. Secreted cytokines and chemokines were effective to stimulate or inhibit cell growth, cell division or apoptosis and cell fusion. Nanometer size particles proved more active in secretion of proinflammatory cytokines than  $\mu$ m size particles.

An option to pharmacologically treat failing implants could be the inhibition of proinflammatory cytokines or chemokines. Acknowledgements

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			Receptor and	Nr. of tested materials
Nr.	Gene Symbol	Gene name	signalling interactions	influencing gene expression
1	Csf1	M-CSF	Cytokine	11
2	Nos2	iNOS	Oxidative Stress	10
3	Arhgap1	Rho GTPase (ac	IL-1 pathway	10
4	Csf3	G-CSF	Cytokine	10
5	Cr2	CD35 (CR2)	Complement	10
6	Dnm1	Dynamin 1	Complement	10
7	Mrc1	CD206	Phagocytose	9
8	ll12b	IL-12(p40)	Cytokine	9
9	Arhgap4	Rho GTPase (ac	IL-1 pathway	8
10	Arhgap26	Rho GTPase (ad	IL-1 pathway	7
11	li10	IL-10	Cytokine	7
12	ll1r1	IL1-Receptor	IL-1 pathway	6
13	Uchl1	Ubiquitin	IL-1 pathway	6
14	Ccl3	MIP-1a	Cytokine	6
15	ll1b	ll-1b	Cytokine	6
16	Fcgr3a	CD16(FCgRIII)	Phagocytose	5
17	Msr1	SR-A (macropha	Phagocytose	5
18	Ctsh	Cathepsin H	IL-1 pathway	5
19	Tnfrsf1b	TNF Receptor 1	IL-1 pathway	5
20	Traf6	TRAF6	IL-1 pathway	5
21	Cd68	CD68	Fc-Receptors	5
22	ll15	IL15	Cytokine	5
23	Dnm2	Dynamin 2	Complement	5
24	Bax	Bax	Apoptose	5
25	Casp9	Caspase9	Apoptose	5

26	Cd14	CD14	Phagocytose	4
27	Hprt1	HPRT	Housekeeping genes	4
28	Ccl2	MCP-1	Cytokine	3
29	Csf2	GM-CSF	Cytokine	3
30	Fcgr1	CD64 (FCgRI)	Complement	3
31	Hip1	HIP1	Complement	3
32	TIr4	TLR4	IL-1 pathway	2
33	Tnfrsf1a	TNF Receptor 1/	IL-1 pathway	2
34	116	II-6	Cytokine	2
35	Tnf	TNF alpha	Cytokine	2
36	Mst1r	CDw136	Phagocytose	1
37	Tnfrsf11a	TNF Receptor 1'	IL-1 pathway	1
38	Fcgr2b	CD32 (FCgRII)	Fc-Receptors	1
39	Siglece	CD170	Complement	1
40	Ms4a2	(FCeRI)	Phagocytose	0
41	Irak1	IL-1-receptor ass	IL-1 pathway	0
42	Nfkb1	NF kappa B1	IL-1 pathway	0
43	Ptges2	Prostaglandin E	IL-1 pathway	0
44	Cdk2	CDK2 (cyclin-de	Housekeeping genes	0
45	Hc	C5a	Fc-Receptors	0
46	Bcl2	Bcl2	Apoptose	0
47	Casp3	Caspase3	Apoptose	0

			Receptor and	up-regulat	down-regu	up-regulat	down-regu	up-regulat	down-reg	unterschie
	Gene Symbol	Gene name	signalling interactions	24h		72h		Differenz	24h-72h	Gesamt-
32	Csf1	M-CSF	Cytokine	4	0	0	7	4	-7	11
45	Nos2	iNOS	Oxidative Stress	5	0	0	5	5	-5	10
41	Arhgap1	Rho GTPase (activating prot. 1)	IL-1 pathway	4	0	0	6	4	-6	10
26	Csf3	G-CSF	Cytokine	5	1	0	6	5	-5	10
33	Cr2	CD35 (CR2)	Complement	3	0	0	7	3	-7	10
6	Dnm1	Dynamin 1	Complement	6	0	0	4	6	-4	10
31	Mrc1	CD206	Phagocytose	4	0	0	5	4	-5	9
24	ll12b	IL-12(p40)	Cytokine	4	0	0	5	4	-5	9
7	Arhgap4	Rho GTPase (activating prot. 4)	IL-1 pathway	2	0	0	6	2	-6	8
12	Arhgap26	Rho GTPase (activating prot. 26)	IL-1 pathway	2	0	0	5	2	-5	7
11	18S	18S	Housekeeping genes	6	0	2	3	4	-3	7
4	li10	IL-10	Cytokine	3	0	0	4	3	-4	7
39	ll1r1	IL1-Receptor	IL-1 pathway	2	3	0	7	2	-4	6
23	Uchl1	Ubiquitin	IL-1 pathway	1	0	0	5	1	-5	6
20	Ccl3	MIP-1a	Cytokine	7	0	1	0	6	0	6
42	ll1b	ll-1b	Cytokine	2	1	0	5	2	-4	6
29	Fcgr3a	CD16(FCgRIII)	Phagocytose	1	0	0	4	1	-4	5
35	Msr1	SR-A (macrophage scavenger r I)	Phagocytose	0	0	5	0	-5	0	5
38	Ctsh	Cathepsin H	IL-1 pathway	0	1	0	6	0	-5	5
34	Tnfrsf1b	TNF Receptor 1B (Receptor 2)	IL-1 pathway	6	0	1	0	5	0	5
36	Traf6	TRAF6	IL-1 pathway	0	0	0	5	0	-5	5
28	Cd68	CD68	Fc-Receptors	0	0	5	0	-5	0	5
47	ll15	IL15	Cytokine	1	0	0	4	1	-4	5
17	Dnm2	Dynamin 2	Complement	5	0	0	0	5	0	5
37	Bax	Bax	Apoptose	0	0	5	0	-5	0	5
21	Casp9	Caspase9	Apoptose	0	0	0	5	0	-5	5

			Receptor and	Differentially
Nr.	Gene Symbol	Gene name	signalling interactions	regulated genes
1	Csf1	M-CSF	Cytokine	11
2	Nos2	iNOS	Oxidative Stress	10
3	Arhgap1	Rho GTPase (ac	IL-1 pathway	10
4	Csf3	G-CSF	Cytokine	10
5	Cr2	CD35 (CR2)	Complement	10
6	Dnm1	Dynamin 1	Complement	10
7	Mrc1	CD206	Phagocytose	9
8	ll12b	IL-12(p40)	Cytokine	9
9	Arhgap4	Rho GTPase (ac	IL-1 pathway	8
10	Arhgap26	Rho GTPase (ac	IL-1 pathway	7
11	18S	18S	Housekeeping genes	7
12	ll10	IL-10	Cytokine	7
13	ll1r1	IL1-Receptor	IL-1 pathway	6
14	Uchl1	Ubiquitin	IL-1 pathway	6
15	Ccl3	MIP-1a	Cytokine	6
16	ll1b	ll-1b	Cytokine	6
17	Fcgr3a	CD16(FCgRIII)	Phagocytose	5
18	Msr1	SR-A (macropha	Phagocytose	5
19	Ctsh	Cathepsin H	IL-1 pathway	5
20	Tnfrsf1b	TNF Receptor 1	IL-1 pathway	5
21	Traf6	TRAF6	IL-1 pathway	5
22	Cd68	CD68	Fc-Receptors	5
23	ll15	IL15	Cytokine	5
24	Dnm2	Dynamin 2	Complement	5
25	Bax	Bax	Apoptose	5
26	Casp9	Caspase9	Apoptose	5
27	Cd14	CD14	Phagocytose	4
28	Hprt1	HPRT	Housekeeping genes	4
29	Ccl2	MCP-1	Cytokine	3
30	Csf2	GM-CSF	Cytokine	3
31	Fcgr1	CD64 (FCgRI)	Complement	3
32	Hip1	HIP1	Complement	3

Nr.	Function	Genome	Genename
1	Housekeeping genes	CDK2 (cyclin-dependent kinase 2)	CDK2
2	Housekeeping genes	GAPDH	Gapdh
3	Housekeeping genes	HPRT	HPRT1
4	Apoptose	Bax	BAX
5	Apoptose	Bcl2	BCL2
6	Apoptose	Caspase3	CASP3
7	Apoptose	Caspase9	CASP9
8	Oxidativer Stress	iNOS	iNOS
9	Phagocytose	CDw136	MST1R
10	Phagocytose	CD206	MRC1
11	Phagocytose	(FCeRI)	MS4A2
12	Phagocytose	CD14	CD14
13	Phagocytose	CD16(FCgRIII)	FCGR3A
14	Fc-Rezeptoren	C5a	НС
15	Fc-Rezeptoren	CD68	Cd68
16	Fc-Rezeptoren	CD32 (FCgRII)	FCGR2B
17	Complement	CD35 (CR2)	Cr2
18	Complement	CD64 (FCgRI)	Fcgr1
19	Complement	CD170	Siglece
20	Complement	SR-A (macrophage scavenger r I)	MSR1
21	Complement	HIP1	HIP1
22	Complement	Dynamin 1	DNM1
23	Complement	Dynamin 2	DNM2

24	Cytokine	G-CSF	CSF3
25	Cytokine	GM-CSF	CSF2
26	Cytokine	IL-10	IL10
27	Cytokine	II-1b	IL1B
28	Cytokine	II-6	IL6
29	Cytokine	IL-12(p40)	IL12p40
30	Cytokine	MCP-1	CCL2
31	Cytokine	Prostaglandin E Synthasse 2	Ptges2
32	Cytokine	TNF alpha	TNF
33	Cytokine	MIP-1a	Ccl3
34	Cytokine	IL15	IL15
35	Cytokine	M-CSF	Csf1
36	IL-1 pathway	IL1-Rezpetor	IL1R1
37	IL-1 pathway	IL-1-receptor associated kinase1	IRAK1
37 38	IL-1 pathway IL-1 pathway	IL-1-receptor associated kinase1 TNF Receptor 1A (receptor 1)	IRAK1 TNFRSF1 A
37 38 39	IL-1 pathway IL-1 pathway IL-1 pathway	IL-1-receptor associated kinase1 TNF Receptor 1A (receptor 1) TNF Receptor 1B (Receptor 2)	IRAK1 TNFRSF1 A TNFRSF1 B
37 38 39 40	IL-1 pathway IL-1 pathway IL-1 pathway IL-1 pathway	IL-1-receptor associated kinase1 TNF Receptor 1A (receptor 1) TNF Receptor 1B (Receptor 2) TRAF6	IRAK1 TNFRSF1 A TNFRSF1 B TRAF6
37 38 39 40 41	IL-1 pathway IL-1 pathway IL-1 pathway IL-1 pathway IL-1 pathway	IL-1-receptor associated kinase1 TNF Receptor 1A (receptor 1) TNF Receptor 1B (Receptor 2) TRAF6 Ubiquitin	IRAK1 TNFRSF1 A TNFRSF1 B TRAF6 UCHL1
37 38 39 40 41 42	IL-1 pathway IL-1 pathway IL-1 pathway IL-1 pathway IL-1 pathway IL-1 pathway	IL-1-receptor associated kinase1         TNF Receptor 1A (receptor 1)         TNF Receptor 1B (Receptor 2)         TRAF6         Ubiquitin         Rho GTPase (activating prot. 1)	IRAK1 TNFRSF1 A TNFRSF1 B TRAF6 UCHL1 ARHGAP1
37 38 39 40 41 42 43	IL-1 pathway IL-1 pathway IL-1 pathway IL-1 pathway IL-1 pathway IL-1 pathway IL-1 pathway IL-1 pathway	IL-1-receptor associated kinase1 TNF Receptor 1A (receptor 1) TNF Receptor 1B (Receptor 2) TRAF6 Ubiquitin Rho GTPase (activating prot. 1) Rho GTPase (activating prot. 4)	IRAK1 TNFRSF1 A TNFRSF1 B TRAF6 UCHL1 ARHGAP1 ARHGAP4
37 38 39 40 41 42 43 44	IL-1 pathway	IL-1-receptor associated kinase1         TNF Receptor 1A (receptor 1)         TNF Receptor 1B (Receptor 2)         TRAF6         Ubiquitin         Rho GTPase (activating prot. 1)         Rho GTPase (activating prot. 26)	IRAK1 TNFRSF1 A TNFRSF1 TRAF6 UCHL1 ARHGAP1 ARHGAP4 ARHGAP2 6
37 38 39 40 41 42 43 44 44	IL-1 pathway IL-1 pathway IL-1 pathway IL-1 pathway IL-1 pathway IL-1 pathway IL-1 pathway IL-1 pathway IL-1 pathway	IL-1-receptor associated kinase1         TNF Receptor 1A (receptor 1)         TNF Receptor 1B (Receptor 2)         TRAF6         Ubiquitin         Rho GTPase (activating prot. 1)         Rho GTPase (activating prot. 4)         Rho GTPase (activating prot. 26)         Rho GTPase (activating prot. 29)	IRAK1 TNFRSF1 DNFRSF1 TRAF6 UCHL1 ARHGAP1 ARHGAP2 6 ARHGAP2 9
37 38 39 40 41 42 43 44 45 46	IL-1 pathway IL-1 pathway	IL-1-receptor associated kinase1         TNF Receptor 1A (receptor 1)         TNF Receptor 1B (Receptor 2)         TRAF6         Ubiquitin         Rho GTPase (activating prot. 1)         Rho GTPase (activating prot. 4)         Rho GTPase (activating prot. 26)         Rho GTPase (activating prot. 29)         Cathepsin H	IRAK1 TNFRSF1 TNFRSF1 TRAF6 UCHL1 ARHGAP1 ARHGAP2 6 ARHGAP2 9 CTSH
37 38 39 40 41 42 43 44 45 46 47	IL-1 pathway	IL-1-receptor associated kinase1         TNF Receptor 1A (receptor 1)         TNF Receptor 1B (Receptor 2)         TRAF6         Ubiquitin         Rho GTPase (activating prot. 1)         Rho GTPase (activating prot. 4)         Rho GTPase (activating prot. 26)         Rho GTPase (activating prot. 29)         Cathepsin H         TLR4	IRAK1 TNFRSF1 A TNFRSF1 TRAF6 UCHL1 ARHGAP1 ARHGAP2 6 ARHGAP2 9 CTSH TIr4



modified from D.M.Underhill and A.Ozinsky, Annu. Rev. Immunol. 2002. 20:825-852

## <u>Mikrocorundum:</u> MR 52 (Al<sub>2</sub>O<sub>3</sub>): GmbH

Manufacturer: particles made according to Bayer-Verfahren (alkalischer Aufschluss von Bauxit, Gewinnung von Al(OH)<sub>3</sub>, Kalzinierung). Primary size1,5 µm (99,8 % Al<sub>2</sub>O<sub>3</sub>;  $\leq$  0,1 % Na<sub>2</sub>O;  $\leq$  0,05 % CaO;  $\leq$  0,04% Fe<sub>2</sub>O<sub>3</sub>;  $\leq$  0,07 % SiO<sub>2</sub>)

Hersteller und Lieferant: Martinswerk

SEM MR52 I. Dörfel (BAM)

M. Gemeinert (BAM): diameter of primary particles around 1.5  $\mu$ m (SEM), specific surface 5,94 m<sup>2</sup>/g (BET-technique), density 3.94 g/cm<sup>3</sup>.

NanocorundumLA  $(Al_2O_3)$ : Auer-Remy VertriebsgesellschaftSeltene Erden, H. Blum & Co. GmbH; Manufacturer: Nanostructured &Amorphous Materials Inc. (USA). Primary size 27 - 43 nm, specific surface35 m²/g, density 3,97 g/cm³. (99,5%  $\alpha$ -Al2O3; 0,3 % Fe2O3; 0,001 % MgO;0,05 % SiO2)



TEM; Nanocorundum (LA) I. Dörfel (BAM) EDX-Spektrum; Nanocorund (LA) I. Dörfel (BAM)

M. Gemeinert (BAM): diameter of primary particles < 100 nm (TEM) specific surface 36 m<sup>2</sup>/g (BET-technique), density 3.677 g/cm<sup>3</sup> EDX-spectrum aluminia ( copper from the nets)





micro corundum sample MR 52



micro graphite sample MX 51





nanocorundum sample NAM



nanographite Printex 90C

nano-chromium-III-oxide