Regulation of Gene Expression by β-Glucans

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Introduction

β-D-glucans are diverse polysaccharides derived from plant cell walls, fungi and bacteria composed of Dglucose monomers linked by (1-3) β -glycosidic bonds and variable amounts of (1-6) and (1-4) branches. β -D-glucans are considered to be "biological response modifiers" because they exhibit antibacterial, antiviral, anticoagulatory, anti-tumoral immunomodulatory and woundhealing activities (Bohn and BeMiller, 1995). In addition, β-D-glucans stimulate the immune system against infectious pathogens and cancer (Saraswat-Ohri et al., 2011; Chan et al., 2009; Aleem, 2013). When ingested in plant materials, β -glucans are absorbed in the small intestine, internalized and fragmented by macrophages and subsequently carried to the spleen, lymph nodes and bone marrow (Chan et al., 2009). Here it is believed that the soluble β -glucan fragments are released and taken up by circulating granulocytes, monocytes and dendritic cells while insoluble fragments induce a cellular response by binding to cell surface receptors. Several types of β -glucan receptors have been identified including several immune receptors, e.g., Dectin-1, Complement Receptor (CR3), CD11b/CD18, Mac-1, aMb integrin (Ross, 2000), Scavenger Receptors (SR),

Abstract: β -D-glucans are diverse polysaccharides derived from plant cell walls, fungi and bacteria. β -D-glucans are composed of D-glucose monomers linked by (1-3) β -glycosidic bonds with varying amounts of (1-6) or (1-4) linked side chains that bind cell surface receptors ultimately triggering changes in intracellular pathways and gene transcription. β -Dglucans have anti-viral, immunomodulatory and anti-cancer activities. This article reviews the current identity and putative function of genes regulated by β -glucans purified from various sources in human cancer and immune cells and in murine dendritic, macrophage cells and lungs. β -D-glucans increase the expression of numerous cytokines and immunomodulatory factors and their receptors in dendritic cells. Pathways and transcription factors involved in the cellular responses to β -glucans are summarized.

Keywords: Glucan, Transcription, Cytokines

Lactosylceramide (LacCer) and toll-like receptors, e.g., TLR-2/6 and trigger responses in macrophages, neutrophils, monocytes, natural killer cells and dendritic cells *in vitro* (Chan *et al.*, 2009; Kim *et al.*, 2011; Legentil *et al.*, 2015). β -glucans also bind to L-Ficolin (Legentil *et al.*, 2015). Through this interaction with plasma membrane receptors, β -glucans potentiate intracellular signaling pathways that ultimately activate transcription factors such as NF κ B resulting in both innate and adaptive immune responses (Li *et al.*, 2010).

In addition to immunomodulatory activity, β -glucans have been reported to have direct anti-cancer activity *in vitro*. A water-soluble β -glucan extract from the mycelia of *Poriacocos* inhibited the viability (MTT assay) of MCF-7 human breast cancer cells with an IC₅₀ of 400 µg mL⁻¹ and decreased cyclin D1 and cyclin E protein expression (Zhang *et al.*, 2006). We reported that a purified preparation of β -D-glucan (from barley) dissolved in DMSO - but not water- inhibited the growth of estrogen receptor α (ER α)+ MCF-7 cells with an IC₅₀ of ~164±12 µg mL⁻¹ compared to normal breast epithelial (ER α -) MCF-10A cells, IC₅₀~464 µg mL⁻¹ (Jafaar *et al.*, 2014). β -glucan inhibited the estradiol (E₂)independent, tamoxifen (TAM) and fulvestrant-resistant, ER α + LCC9 and LY2 cells (derived from MCF-7 cells)



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with IC₅₀ values of 4.6±0.3 and 24.2±1.4 µg mL⁻¹, respectively (Jafaar *et al.*, 2014). In contrast, the "triple negative/basal-like" MDA-MB-231 breast cancer cells were not growth inhibited by β -glucan (Jafaar *et al.*, 2014). Conversely, aqueous-soluble β -glucans reportedly had no direct cytotoxic effects on a panel of common cancer cell lines including blastoma, carcinoma and sarcoma cells (Chan *et al.*, 2009).

There are many studies of β -glucans purified from the cell walls of yeast, fungi and plants as immunomodulators that increase the levels of proinflammatory cytokines, chemokines and cell adhesion molecules (Novak and Vetvicka, 2008). We will review immunomodulatory genes regulated by β -glucans in human and mouse cells. In addition, we reviewed the literature to identify additional genes increased or inhibited by glucan treatment of cells in vitro. We downstream effects of these summarize the transcriptional changes and how they inform the mechanisms by which glucans act in various cell types.

Glucans Increase Gene Transcription in Human Cancer Cells and Immortalized Normal Cells

A number of reports have examined the effect of β glucan on transcription in human and murine immune cells with a few studies examining the activity of glucans in human cancer cells. Table 1 summarizes those genes that were transcriptionally induced by β glucan in the cells examined. First, we will review the results in human cells.

β -Glucan in Breast Cancer Cells

In our lab, it was shown that a purified β -D-glucan isolated from barley and dissolved in DMSO, but not water, inhibited cell viability in two human breast cancer cells lines: MCF-7 and LCC9, an endocrine-resistant (estradiol, tamoxifen and fulvestrant) cell line derived from MCF-7. Using two concentrations of β -D-glucan, we examined the expression of a set of genes implicated in breast cancer in both cell lines using PCR array analysis.

Table 1. Genes upregulated by glucan in cell systems. The concentration, form and source of β -glucan used in the experiment is indicated. The comments include information about the genes regulated taken from the references cited and the GeneCards human gene database http://www.genecards.org/I.MW = low molecular weight

Gene (s)	Cell system	β-glucan	Ref.	Comments
BIRC5, BRCA1, BRCA2, CCNA1, PGR, RASSF1	MCF-7 human breast cancer cells	10 or 50 μ g mL ⁻¹ β -D glucan purified from barley and purchased from Sigma (cat. No. G6513, purity ~97%), dissolved in DMSO	(Jafaar <i>et al.</i> , 2014)	BIRC5 is anti-apoptotic;BRCA1 and BRCA2 are involved in DNA repair, Cyclin A1(<i>CCNA1</i>) regulates cell cycle progression, progesterone receptor (<i>PGR</i>) is a breast cancer differentiation marker, RASSF1 (<i>NORE2A</i>) is a scaffolding protein that functions as a tumor suppressor
CTSD, PTGS2	MCF-7 human breast cancer cells	$10 \ \mu g \ m L^{-1} \ \beta$ -D glucan	(Jafaar <i>et al.</i> , 2014)	Cathepsin D (<i>CTSD</i>) is an acid protease; COX-2 (<i>PTGS2</i>) increases pro-inflammatory cytokines
MK167	MCF-7 human breast cancer cells	50 μg mL ^{-1} β-D glucan	(Jafaar <i>et al.</i> , 2014)	Ki-67 is a marker of cell proliferation
EGF, GL11, HIC, IGF1, IGFBP3, PTGS2	LCC9 endocrine-resistant human breast cancer cells	10 or 50 μ g mL ⁻¹ β -D glucan	(Jafaar <i>et al.</i> , 2014)	GLI1 is a zinc finger transcription factor; HIC is the gene <i>MDFIC</i> is a transcriptional regulator
TP53, CDKN1B	MCF-7 human breast cancer cells	Fungal β -glucan (Botryosphaeran, BOT) was produced by <i>B. rhodina</i> MAMB-05 grown on nutrient medium containing glucose (BOT _{GLC}) and fructose (BOT _{FRU} Cells were treated with 100 µg mL ⁻¹ for 48 h	(Queiroz <i>et al.</i> , 2015)).	P53 and P27 are involved in cell cycle arrest
CD86			(Harnack <i>et al.</i> , 2011)	Maturation marker-measured cell surface expression by flow cytometry. Observed peak increase at 48 h tx. Both cell lines express dectin-1, the major β-glucan receptor.
HSP70 protein, MUC1	Human gastric carcinoma cells	LMW β -glucan purified from conditioned medium of <i>Aureobasidium pullulans</i> GM- NH-1A1 with ~ 70% β -(1-6) branches 100 μ g mL ⁻¹	(Tanaka <i>et al.</i> , 2011)	MUC1 is mucin 1, a transmembrane protein.

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<i>PKC, CXCL8</i> (IL-8 gene), <i>CDC42, BBC3</i>	Detroit-573 human fibroblasts and Ha	6 different synthetic glucans- no concentrations or treatment	(Vetvicka et al., 2011)	Regulators of cell growth and PUMA is a BCL-2 pro-
(PUMA)	Cat human keratinocytes	times indicated		apoptotic protein
L-10, IL-12, IL-23 ranscripts	Human monocyte-derived Dendritic Cells (DCs) from healthy donors	zymosan (β-glucan from Saccharomyces cerevisiae (InvivoGen) 10 and $200 \ \mu g \ mL^{-1}$	(Gerosa et al., 2008)	reviewed in (Lyakh <i>et al.</i> , 2008)
L23A, PTGS2	Human monocyte-derived DCs from healthy donors	$\beta(1,3)$ -glucan Zymosan from Saccharomyces cerevisiae (Sigma) 1 mg mL ⁻¹ ,4 h	(Rodríguez et al., 2014)	β-glucan activates the Unfolded Protein Response in DCs
TGF-a, IFN-r, IFN-a,, IL-29, IL-28B, IL-28A, IL-7, IL-23A, IL-19, IL-15, IL-12B, IL-12A, IL10, IL8, IL7, IL-6, IL-1B, IL-1A, EB13, IL22RA1, IL15RA, IL7R, IL6R, IL4R, IL3RA, IL2R: CCR7, CCR6, CX3CL1, CXCL11, CXCL10, CXCL CXCL2, CXCL3, CCL3, CXCL2, CXCL1, CCL22, CCL20, CCL19, CCL17, CCL15, CCL4, CCL3, CCL1, LILRB4, LAMP3, ECGF1, CD86, CD83, CD80, CD58, CD54, CD47, CD44, ADAMDEC ISG20, IFITM3, IFIT4, IFIT2, IFIT1, IFI44, IFI35, IF127, GIP3, GIP2 RELA, RELB, NFKB2, NFKB1, EB13, CFLAR, BTG1, BASP1, ATF4, AIL20, CD80, CD54, CAPA	DCs from healthy donors 4, .9,	PS-G, a branched (1→6)-β- d-glucan) purified from <i>Ganoderma lucidum</i> 10 µg mL ⁻¹ for 16 h	(Lin <i>et al.</i> , 2006)	AffymetrixGeneChip microarray
NFATCA1, IFNB1, S100A9, S100A8, GAPDH, TGFA, MMP2, CYP7B1, IL12B, IL12A, EB13, ITGB8, IL27, IL15, CD38, CD80, CCR7, STAT3, IRF1, TNF, CCL4, IL10,IL20, IL23, NFKB1A,NFKBIZ, PTGS2, TLR2, SOD2,	DCs from healthy donors	β-glucan from baker's yeast (Sigma-Aldrich) 10 μg mL ⁻¹ 4 or 12 h	(Cardone <i>et al.</i> , 2014)	Agilent-014850 Whole Human Genome Microarray 4×44 K G4112F GEO accession number GSE42189.
NFKB1, IL6, IL1B, IL9 INF-α, IL6, IL8, IL12, IL18	Human B lymphocytes isolated from healthy donors	Curdlan (Sigma-Aldrich) 100 ug mL^{-1} for 24 h	(Ali <i>et al.</i> , 2015)	IL-8 expression was shown to involve Dectin-1, SYK, MAPKs and transcription factors AP-1 and NF-kB
GITRL protein	Murine DCs	whole β glucan particles from <i>S. cerevisiae</i> treated for 48 h (100 μ g mL ⁻¹)	(Tian et al., 2012)	<i>TNFSF18</i> is the gene for GITRL a cytokine in the TNF ligand family
<i>Ddx58, Mda5,</i> IL-1β, L-6, TNFα, GM-CSF nd G-CSF	Murine macrophage- derived RAW264.7 cells	Aureobasidium pullulans produced purified β (1->3), (1->6) D glucan 100 µg mL ⁻¹) time course experiments: 3-24 h	(Muramatsu <i>et al.</i> , 2012)	<i>Ddx58</i> is RIG-1 (retinoic acid- inducible gene-I). MDA5 is an intracellular pattern recognition molecule
<i>UMPS</i> =Uridine Monophosphate synthetase,	Human KB cells (subclone of HeLa cells) grown as a tumor xenograft in female athymic mice	<i>Lentinus edodes</i> (Shiitake mushroom) Dose (ip injection): 0.1 mg/kg/day 2X/week for	(Harada <i>et al.</i> , 2010)	measured by RT-PCR in the microdissected tumor; UMPS is also called Orotate Phosphoribosyl Transferase (OPRT)
IFN-Y	Lungs of normal mice	3 weeks Antrodia camphorate beta- glucan; mice were fed beta glucan daily for 12 days	(Wang et al., 2015)	(OPRT)

 β -D-glucan increased the expression of 9 genes in MCF-7 human breast cancer cells and 5 genes in LCC9 breast cancer cells (Jafaar et al., 2014). These genes are summarized in Table 1. Overall, depending on the concentration of β -D-glucan, we observed that β -Dglucan increased the expression of genes involved in proliferation (MK167 (Ki-67), CCNA1 (cyclin A1) and BIRC5), invasion (CTSD (cathepsin D)), inflammation (PTGS2 (COX-2)), differentiation (PGR (progesterone receptor)), tumor suppression (RASSF1) and DNA repair (BRCA1, BRCA2). Surprisingly, a set of different genes were upregulated by β -D-glucan in the endocrine-resistant (tamoxifen and fulvestrant) LCC9 cells with the exception of the common increase in PTGS2. These genes were involved in proliferation (EGF, IGF1, IGFBP3) and transcription (GL1). The mechanism for this difference is unknown, although LCC9 cells have higher NFkB expression/activity than MCF-7 (Litchfield et al., 2014).

More recent studies reported that two preparations of fungal β -glucans inhibited MCF-7 cell viability, stimulated apoptosis and ROS production and increased necrosis (Queiroz *et al.*, 2015). BOT_{GLC} and BOT_{FRU} (100 µg mL⁻¹, Table 1) increased the mRNA and protein expression of p53 (*TP53*) and p27 (*CDKN1B*) after 48 h of treatment (Queiroz *et al.*, 2015). The authors reported that BOT_{GLC} activated AMPK, increased FOXO3a protein and reduced phospho-FOXO3a, suggesting that stimulation of FOXO3a transcriptional activity is the mechanism by which cell arrest is achieved. This was supported with the observation that treatment with these β -glucans resulted in an increase in p53 and p27 transcription (Queiroz *et al.*, 2015).

β-Glucan Regulates Protein and Gene Expression in Gastric Cancer Cells and other Human Cell Lines

Treatment of human gastric carcinoma cells with β glucan increased the expression of HSP70 protein (*HSPA1A* gene) and *MUC1* transcript levels (mucin-1, a transmembrane protein) - both of which are protective factors in the gastric mucosa (Tanaka *et al.*, 2011). We note that the primary focus of this study was to examine the protective effects of β -glucans on gastric lesions in mice with confirmation of HSP70 and MUC1 in gastric cancer cells.

Synthetic glucans increased expression of *PKC* (protein kinase C), *CXCL8* (IL-8), CDC-42 and PUMA (*BBC3*) transcripts in human Detroit-573 fibroblasts and HaCaT keratinocytes as measured by non-quantitative PCR (Vetvicka *et al.*, 2011). These genes are regulators of cell growth and PUMA is a BCL-2 pro-apoptotic protein.

Microarray Analysis in Human Monocyte-Derived Dendritic Cells

Gene array profiling of human monocyte-derived Dendritic Cells (DCs) treated with a purified extract of Polysaccharide (PS-G, a branched $(1\rightarrow 6)$ - β -d-glucan) from Ganoderma lucidum, a Chinese medicinal mushroom, identified 3477 (17%) probe sets upregulated (2-fold) and 4418 (19%) probe sets downregulated (2fold) after 16 h of treatment (Lin et al., 2006). The identity of genes included in the text of that manuscript (Lin et al., 2006) are shown on Table 1 and 2. Significant increases in transcript levels were observed for a number of cytokines (IL-12A, IL-12B, IL-23A, IL-27 and EBI2; also known as IL-27B), chemokines and chemokine receptors (CCL20, CCL19, CCL5, CXCL10, CXCL11 and CCR7) and cell surface proteins (CD80, CD83 and CD86). The authors also noted that an increase in transcripts for NFKB1, NBKB2, RELA, RELB and MAPK11 which support the model that β -glucans activate human dendritic cells through the NFkB and p38 MAPK pathways. Further, the authors suggested that the increase in CCL20, IL-27, IL-23A, IL-12A and IL-12B transcripts after PS-G treatment of human DCs may play a role in the anti-tumor activity of β-glucans (Lin et al., 2006).

A more recent microarray profiling of human monocyte-derived DCs treated with β -D glucan (10 μ g mL^{-1}) for 4 or 12 h identified ~1500 genes that were either inhibited (38%) or induced (62%) (Cardone et al., 2014). The β-D glucan-regulated genes were divided into 6 groups: (1) Early-inhibited genes (4-6 h after activation); (2) genes inhibited similarly at both early and late times (early-late-inhibited genes); (3) lateinhibited genes; (4) early-induced genes; (5) genes induced similarly at both early and late times (earlylate-induced genes; and (6) late-induced genes (12 h after activation). Gene ontology analysis indicated that the early induced genes were likely involved in apoptosis or stress and antiviral responses, e.g., IFNB1, NFKBID, NFATC1, TNF, proIL1β, NFKBIZ and that these genes were potentially regulated by Interferon Regulatory Factors (IRFs). The early-induced genes were identified to be involved in chemotaxis, IL-1 production and NF κ B or STAT signaling and were predicted to be regulated by NFkB and IRFs. The lateinduced genes included chemokines, cytokines and factors involved in DC activation other and proliferation, e.g., IL6, 110, IL12B, IL20, IL23A, IL1F9, CSF7, CSF3; and were predicted to be regulated by NFkB, IRFs, AP1, STATs and CEBPs. The authors demonstrated that β -glucan stimulates IL-1 expression which once processed by the caspase/inflammasome pathway acts in an autocrine fashion to stimulate the expression of the late-induced cytokines at the transcriptional level via IκB-ζ modulation (Cardone et al., 2014). Further, the authors reported that β -glucan also promoted an IFN-I gene signature consistent with the reported induction of IFN-I in response to fungi via Dectin-1 signaling. These data demonstrated that IL-1 and IFN- γ differentially regulate the β -glucan-induced Th responses in human DCs.

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Table 2. Genes down-regulated by glucan. The cell type	and concentration and form of β-gluc	an used in these studies is indicated. The c	comments include
information about the genes regulated taken from	n the references cited and from the Ge	ene Cards human gene database http://www	v.genecards.org/

Gene	Cell system	β-glucan	Ref.	comments
Measured Cyclin D1 and Cyclin E proteins by flow cytometry	MCF-7 human breast cancer cells	$400 \ \mu g \ mL^{-1} \ PCM3-II$, a water- soluble β -glucan purified from the mycelia of <i>Poriacocos</i> , 48 h treatment	(Zhang et al., 2006)	Cell cycle regulators
CDKNIC, PLAU, RARB	MCF-7 human breast cancer cells	10, 50 μg mL ⁻¹ β-D glucan; 24 h	(Jafaar <i>et al.</i> , 2014)	CDKN1C is a negative regulator of cell proliferation; Plasminogen Activator, Urokinase (<i>PLAU</i>) is a serine protease, Retinoic Acid Receptor β (<i>RARB</i>) is a differentiation maker
CTNNB1, IGFBP2, SLIT2	MCF-7 human breast cancer cells	50 µg mL ⁻¹ β -D glucan; 24 h	(Jafaar <i>et al.</i> , 2014)	<i>CTNNB1</i> is β catenin; <i>IGFBP2</i> is Insulin Like Growth Factor Binding Protein 2; SLIT2 is a secreted glycoprotein involved in cell migration
MUC1, SNAI2	MCF-7 human breast cancer cells	10 μg mL ⁻¹ β-D glucan; 24 h	(Jafaar <i>et al.</i> , 2014)	MUC1 is a transmembrane protein that is cleaved at its intracellular domain interacts with ERα and contributed to endocrine-resistance (Kufe, 2013); <i>SNA12</i> is the protein LUG which is a transcriptional repressor
ADAM23, BRCA2, CDH13, CDKN1C, CTNNB1	LCC9 endocrine-resistant human breast cancer cells	10, 50 μg mL ⁻¹ β-D glucan; 24 h	(Jafaar <i>et al.</i> , 2014)	ADAM23 is a membrane protein that acts as a tumor suppressor; <i>CDH1</i> is Cadherin 13; <i>CDH13 is</i> Cyclin-Dependent Kinase Inhibitor 1C- a negative regulator of cell proliferation; <i>CTNNB1</i> is β-catenin
ERCC5	HepG2 cells	A β-glucan-containing polysaccharide complex purified from <i>Agaricusblazei</i> Murill mushroom of Brazilian origin by aqueous extraction (Gonzaga <i>et al.</i> , 2005) 50 μ g mL ⁻¹ for 6 h	(da Silva <i>et al.</i> , 2013)	ERCC5 is involved in excision repair following DNA damage
Identified 61 downregulated miRNAs -focused on <i>mmu-miR-9</i> <i>CREB mRNA and</i> protein	ç	100 ug mL ^{-1} for 24 h; whole β -glucan particles from S. <i>Cerevisiae</i>	(Tian <i>et al.</i> , 2015)	miRNA microarray assay GEO accession number GSE67578; Identified Runx1, a transcription factor involved in differentiation of MDSCs, as a direct, <i>bona fide</i> target of miR-9. CREB, transcription factor, suppressed miR-9 transcription.
TS, DPD		e lentinan purified from <i>Lentinus edodes</i> (Shiitake mushroom) Dose (ip injection): 0.1 mg/kg/day 2X/week for 3 weeks	(Harada <i>et al.</i> , 2010)	measured by microdissection and RT-PCR; Thymidylate Synthase (TS) Dihydropyrimidine Dehydrogenase (DPD)
ARHGDIB, ATM, CEBPA, CSFIR, CST, F13A1, ITGAM, ITGB2, VCL	Human monocyte-derived DCs from healthy donors	PS-G, a branched $(1\rightarrow 6)$ -β- D-glucan) purified from <i>Ganoderma lucidum</i> 10 µg/ml for 16 h	(Lin et al., 2006)	Affymetrix GeneChip microarray
Ptgs2 (Cox2 gene)	Lungs of normal C57BL/6JNarl mice	Antrodia camphorate beta- glucan; mice were fed beta glucan daily for 12 days	(Wang <i>et al.</i> , 2015)	

β-Glucan Stimulates IL23A and IL-10 Transcription But not IL-12 in Human Dendritic Cells

In a study designed to determine the effect of β -glucan on the production of IL12 and IL23 in human

dendritic cells, high and low concentrations of purified β -glucan (10 *versus* 200 μ g mL⁻¹ zymosan) were shown to induce transcription of IL-23 and IL-10, which was enhanced when cells were simultaneously treated with R848 (a Toll-like receptor

7 and 8 agonist) or with Pam2C (TLR2 ligand) (Gerosa *et al.*, 2008). IFN- γ priming inhibited zymosan-stimulated IL-23 transcription (Gerosa *et al.*, 2008). In contrast, transcription of IL-12 was not induced by zymosan alone. Instead, expression of IL12 was observed only when cells were first primed with IFN- γ and treated with low concentrations of zymosan and R848. Based on these observations, the authors concluded that the differential expression of IL-23 and IL-12 is dependent on the specific PPRs (pattern recognition receptors) that are activated by different microorganisms. Specifically, TLR2 is able to differentially regulate the expression of these two cytokines which allows DCs to respond appropriately to different pathogens (Gerosa *et al.*, 2008).

In a more recent study, Rodríguez et al. (2014) showed that selective upregulation of IL-23 by mg mL^{-1}) involves zvmosan (1 activating transcription Factor 2 (ATF2) in addition to NFKB which stimulates both IL-23 and IL-10. In these studies, the authors suggest that zymosan interaction with dectin-1 activates PKA-MEK-ERK and PKC signaling leading to activation of ATF2 and coordinates with TLR2 signaling to activate NFkB. ATF2 and NFκB stimulate IL23A transcription (Rodríguez et al., 2014). While zymogen and two other insoluble β -glucans (curdlan- and pustulancontaining latex beads) resulted in the disappearance of the Endoplasmic Reticulum Stress Response (ERS) transcription factors CHOP and XBP-1 from the nucleus of human DCs and macrophages, there was no evidence that CHOP is involved directly in IL-23 induced expression (Rodríguez et al., 2014).

β-Glucans Induce Cytokine Production in Human B Lymphocytes

While most studies have focused on the activation of gene expression in DCs, a recent report demonstrated that purified β -glucan (curdlan) induces expression of several cytokines including TNF α , IL-6 andIL-8 in human B lymphocytes at both the mRNA and protein level (Ali *et al.*, 2015). Activation of IL-8 was shown to be mediated through the Dectin-1 receptor and involved SYK, ERK and JNK and activation of the transcription factors AP-1 and NF κ B. Interestingly, β -glucan had no effect on antibody production or proliferation suggesting that β -glucans induce a different response than TLR-9 agonist CpG oligodeoxynucleotide (CpG) that is a bacterial and viral DNA component known to induce cytokine production in B lymphocytes (Ali *et al.*, 2015).

Glucans Increase gene Transcription in Murine (Mouse) Immune Cells

Treatment of murine Myeloid-Derived Suppressor Cells (MDSCs) with whole β -glucan particles was

shown to increase the expression of Runx1, a transcription factor that induces differentiation in Myeloid-Derived Suppressor Cells (MDSC) and *mmu*-*miR-181d in M-MDSCs (downregulated in G-MDSCs)* (Tian *et al.*, 2015). These authors did not examine the function of miR-181d and a search in PubMed revealed no studies examining miR-181d in MDSCs. miR-181d is a stress-responsive miRNA in thymocytes with expression decreasing in response to LPS (Belkaya and van Oers, 2014). A search of microRNA.org revealed 8084 targets for has-miR-181d, but the role of increased miR-181d in response to β -glucan in MDSCs remains to be defined.

Treatment of murine bone marrow derived DCs with Whole β -Glucan Particles (WGPs) rapidly activated the dectin-1 receptor, increased SYK phosphorylation (10-20 min) and increased cell surface GITRL protein (RNFSF18) levels measured 48 h later (Tian et al., 2012). GIRTL (also called TL6) is a cytokine member of the TNF ligand family. GIRTL primes cytotoxic T responses lymphocyte and downregulates the suppressive activity of regulatory T cells, thus inhibiting tumor progression in the mouse Lewis Lung Carcinoma (LLC) cell model (Tian et al., 2012). The authors also showed that treatment of mice implanted with LLC tumor cells with WBPs resulted in an increase in GITRL on DCs in vivo and slower tumor development.

Time course experiments in murine macrophagederived RAW264.7 cells showed that β (1->3),(1->6)-Dglucan, purified from Aureobasidium pullulan, strongly upregulated the inflammatory cytokines IL-1β and IL-6 and weakly upregulated TNFa. Increases in GM-CSF (CSF2) and G-CSF (CSF3), cytokines important for differentiation and proliferation, were also observed by 3 h with increased Ddx58 (RIG-1- retinoic acid inducible gene-1) and Mda5 (melanoma differentiation-associated protein 5) seen 6 h after treatment (Muramatsu et al., 2012). RIG-1 is necessary for Type I interferon production when cells are infected with a virus and MDA5 is an intracellular pattern recognition molecule for virus-derived RNAs which is necessary for Type I interferon production when cells are virus infected. These results correlate with the anti-viral activity of β glucan observed in mice (Muramatsu et al., 2012).

Treatment of RAW264.7 mouse macrophage cells with β -glucan increased TNF α and GM-CSF (granulocyte macrophage colony stimulating factor) after 3 h. CSF2 is involved in the development and differentiation of lymphocytes from stem cells increased at 3 h and remained upregulated through 24 h.

β-Glucan Regulates Genes in a Human Tumor Cell Xenograft

Treatment of female athymic mice with lentinan (LNT, Glc(b1-3)[Glc(b1-6)]Glc(b1-3)Glc(b1-

3)Glc(b1-3)[Glc(b1-6)]b-Glc,

https://pubchem.ncbi.nlm.nih.gov/compound/37723) purified from Lentinusedodes (Shiitake mushroom) at a dose of 1 mg/kg/day 2X/week for 3 weeks reduced growth of xenografted human KB cells (a subclone of HeLa cells) (Harada et al., 2010). The KB tumors showed reduced expression of UMPS (Uridine monophosphate synthetase, also called Orotate Phosphoribosyl Transferase (OPRT)) at both mRNA and protein levels (Harada et al., 2010). Co-treatment with lentinan and S-1, an oral anticancer drug, increased upregulation of UMPS. Inhibition of UMPS would inhibit pyrimidine biosynthesis and thus inhibit nucleic acid biosynthesis, correlating with the antitumor activity of lentinan (Ina et al., 2013). The authors suggested that since lower levels of expression of OPRT has been associated with resistance to 5-FU based chemotherapies, cotreatment with lentinan may increase the successful use of therapeutic drugs such as S-1 and other 5-FU based chemotherapies.

Genes Inhibited by β -Glucan

A number of studies have identified genes whose expression is decreased by β -glucan treatment in human and mouse cells. Table 2 summarizes the information on these genes that have been identified. First, the results in human cells will be reviewed followed by studies in mouse cells.

β-Glucan Inhibits Gene Transcription in Breast Cancer Cells

Early studies in MCF-7 cells showed that a 48 h treatment with 400 μ g mL⁻¹ PCM3-II, a water-soluble β -glucan, inhibited cell viability and coordinately reduced Cyclin D1 and Cyclin E protein expression (Zhang et al., 2006). However, that study did not examine the impact of PCM3-II on gene transcription. We examined the effect of two concentrations of β-Dglucan on the expression of a set of genes implicated in breast cancer in MCF-7 and LCC9 cells using a PCR array. We reported that β -D-glucan inhibited the expression of cell proliferation and differentiation markers in MCF-7 cells, e.g., CDKN1C, CTNNB1, ERBB2, MUC1 and RARB (Jafaar et al., 2014). These changes in gene expression correspond to the observation that β -D-glucan dissolved in DMSO, but not water, inhibits MCF-7 cell proliferation (Jafaar et al., 2014). β-D-glucan also inhibited the expression of CDKN1C and CTNNB1 in LCC9 cells (Jafaar et al., 2014). The mechanism(s) for these effects in MCF-7 and LCC9 cells has not been examined.

β -Glucan Inhibits ERCC5 Expression in HepG2 Cells

Treatment of HepG2 human hepatoma cells with a β-glucan-containing polysaccharide complex purified from Agaricusblazei (Murill mushroom of Brazilian origin) by aqueous extraction (Gonzaga et al., 2005) had no effect on cell viability (MTT assay), CYP1A1 or CASP9 gene transcript levels, but inhibited ERCC5 expression (da Silva et al., 2013). ERCC5 is involved in excision repair following DNA damage, but the authors did not pursue this finding. Interestingly, βglucan treatment of HepG2 cells increased metabolites involved in bioenergetic metabolism, including alanine, glutamate and creatine. From these studies, the authors concluded that β -glucan increases metabolites needed for high energy metabolism and stimulates bioenergetics (da Silva et al., 2013). injection croaker Interestingly, of yellow (*Pseudosciaenacrocea*) fish with β -glucan (5 mg kg⁻¹ body weight, from Sigma) 6 h prior to hypoxic stress inhibited reactive oxygen formation in the liver and increased transcription of Pyruvate Kinase (PK) while reducing F-ATPase, Succinate Dehydrogenase (SDH) and Malate Dehydrogenase (MDH) expression (Zeng et al., 2016). Which subunit of SDH or form of MDH was examined by PCR was not indicated. These results suggest that β -glucan inhibits the TCA cycle and enhances glycolysis in vivo in the liver of yellow croaker fish, although direct examination of TCA cycle metabolites, oxygen consumption or ATP production was not performed.

β-Glucan Inhibits miR-9 Expression in Mouse Macrophage Cells

A microRNA microarray in Whole β-Glucan Particles (WGP)-treated MDSCs isolated from the spleen of Lewis lung carcinoma bearing mice identified 61 miRNAs that were downregulated whereas 40 miRNAs were upregulated (Tian et al., 2015). Specifically, the authors identified miR-9 as an important regulator of Runx1 (runt-related transcription factor) as a direct, bona fide target of miR-9 (verified by 3'UTR luciferase reporter assay in HEK-293T cells). They also reported a decrease in Creb transcript levels in WGP-treated MDSCs, identified a binding site for CREB in the miR-9 promoter region and verified CREB suppression by a promoter luciferase reporter assay. Both CREB and mi-RNA suppression by WGP were shown to be mediated by the Dectin-1 pathway. These results support the conclusion that through the Dectin-1

receptor, CREB directly regulates miR-9 expression in response to β -glucan. The downregulation of miR-9 results in an increase in Runx1 allows for differentiation, decreased immunosuppressive function and reduced tumor growth (Tian *et al.*, 2015).

β-Glucan Inhibits Proteins that Regulate Nucleic Acid Synthesis in a Human Tumor Cell Xenograft

As reviewed above, treatment of female athymic mice with lentinan at a dose of 1 mg/kg/day 2X/week for 3 weeks reduced xenograft human KB cell tumors (Harada *et al.*, 2010). The authors reported reduced thymidylate synthase and dihydropyrimidine dehydrogenase proteins in the KB xenografts, results that correlated with inhibition of KB xenograft growth *in vivo* (Harada *et al.*, 2010).

As discussed above, gene array profiling of human monocyte-derived DCs treated with a PS-G identified 4418 (19%) probe sets downregulated (2-fold) after 16 h of treatment (Lin *et al.*, 2006). The identity of downregulated genes included in the text of that manuscript (Lin *et al.*, 2006) is shown in Table 2. The authors suggested that the decrease in *ARHGDIB*, *ATM*, *CEBPA*, *CSFIR*, *CST*, *F13A1*, *ITGAM* and *ITGB2* transcripts after PS-G treatment may play a role in the

anti-tumor activity of β -glucans (Lin *et al.*, 2006). Another microarray profiling of human monocytederived DCs treated with β -D glucan (10 µg mL⁻¹) for 4 or 12 h identified ~ 1500 genes that were either inhibited (38%) or induced (62%) (Cardone *et al.*, 2014). The authors focused on upregulated transcripts and none of the downregulated genes were described; thus, these are not included in Table 2.

Conclusion

From the genes identified as regulated by β -glucan in this review, it is clear that β -glucan regulates the transcription of numerous immunomodulatory genes in human and mouse dendritic and macrophage cells. In addition, gene expression changes in response to β glucans have been identified in human breast, gastric and liver cancer cells as well as a few other cell lines as summarized in Table 1 and 2. Figure 1 is a model of the generalized pathway by which β -glucan regulates gene transcription by interaction with a plasma membrane receptor which then activates an intracellular pathways leading to activation or inhibition of transcription factor activity. The activated transcription factors then regulate the transcription of cell-specific target genes.



Fig. 1. General model of β-glucan regulation of gene transcription. Examples of cell type specific receptors, pathways and transcription factors are in plain font

Authors Contributions

MOH and CMK contributed equally to the writing of this review.

Conflict of Interest

The authors declare no conflict of interest.

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