



INTERACCIÓN DE LOS PROBIÓTICOS CON LA MICROBIOTA INTESTINAL Y LA OBESIDAD

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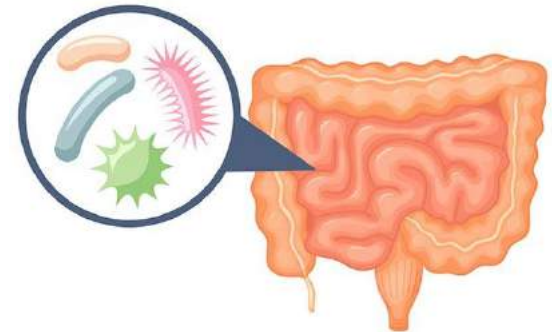
Divulgación de conflicto de intereses

Esta presentación NO es respaldada por ninguna subvención

Probióticos

“Microorganismos vivos que, cuando se administran en cantidades apropiadas, confieren al huésped un beneficio para la salud”

FAO/OMS 2001



Composición de la Microbiota Intestinal

Cinco filos



Bacteroidetes

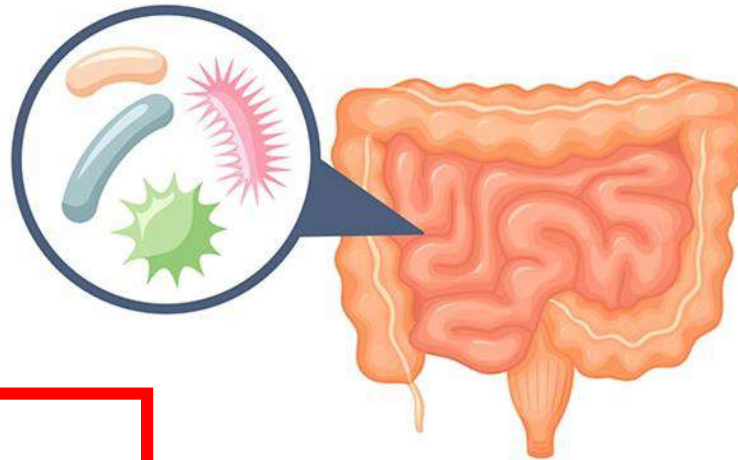
Firmicutes

Actinobacteria

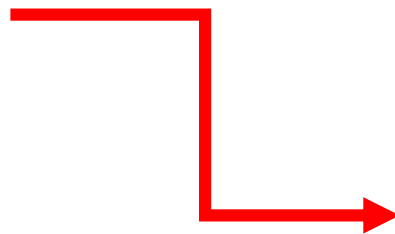
Proteobacteria

Fusobacteria

Verrucomicrobia



baja concentración de oxígeno
más de 800 especies bacterianas
hasta 10¹² células / gramo de heces

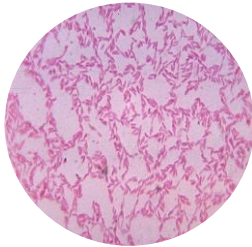


90% de las bacterias totales



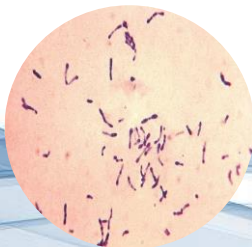
Firmicutes

Los géneros notables incluyen *Bacillus*, *Enterococcus*, *Streptococcus*, *Staphylococcus*, *Listeria*, ***Lactobacillus*** (en la clase Bacilli), *Clostridium* (en la clase Clostridia) y *Mycoplasma* (en la clase Mollicutes).



Bacteroidetes

Una de las especies del género clínicamente más importantes es el *Bacteroides fragilis*.



Actinobacteria

Las especies del género ***Bifidobacterium*** desempeñan un papel fundamental en la regulación de la actividad intestinal normal, principalmente en bebés lactantes.

Especies bacterianas aceptadas como probióticos

Lactobacillus:

Acidophilus

Casei

Fermentum

Gasseri

Johnsonii

Paracasei

Plantarum

Rhamnosus

salivarius

Bifidobacterium:

Adolescentes

Animalis

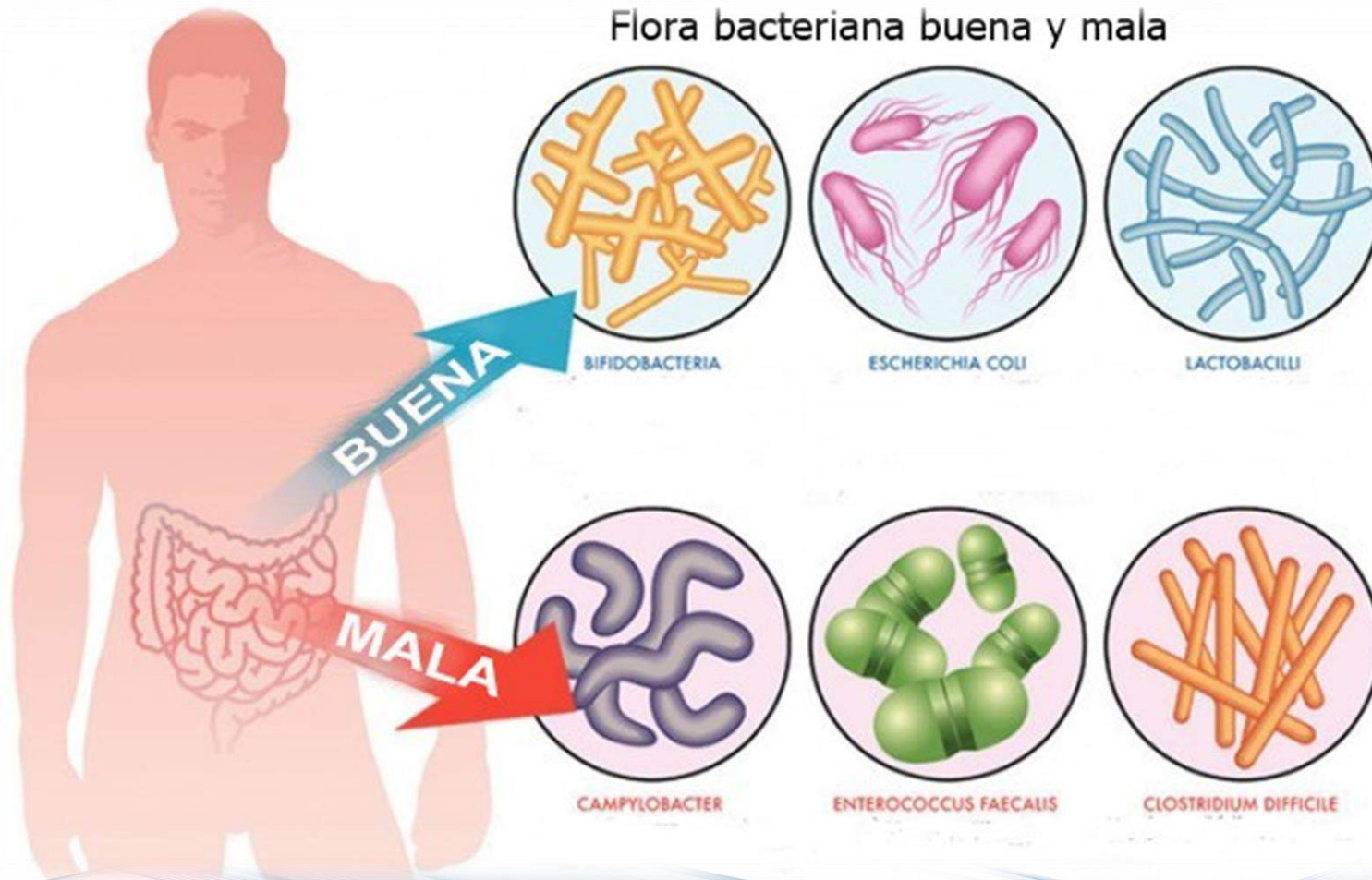
Bifidum

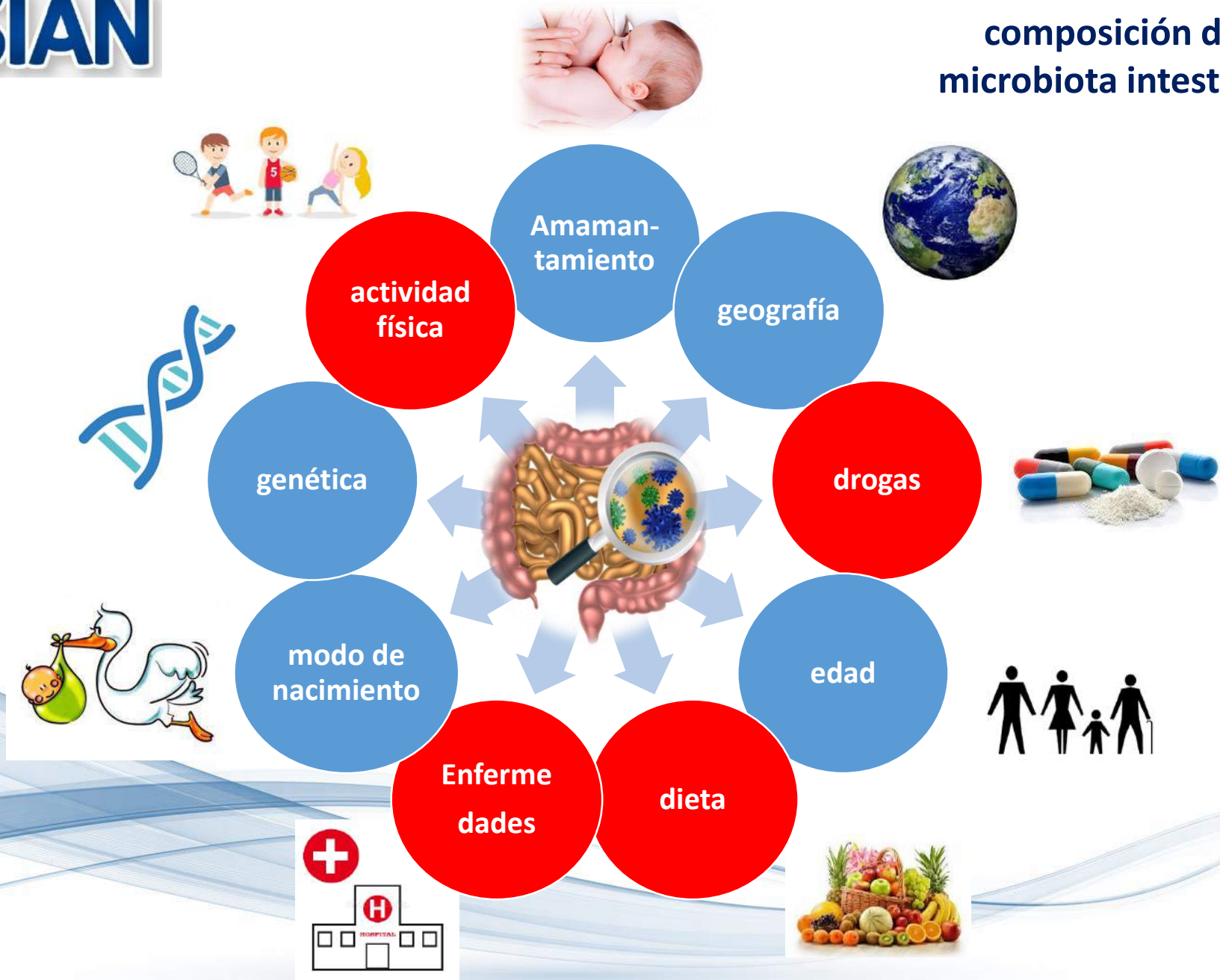
Breve

longum



Flora bacteriana buena y mala



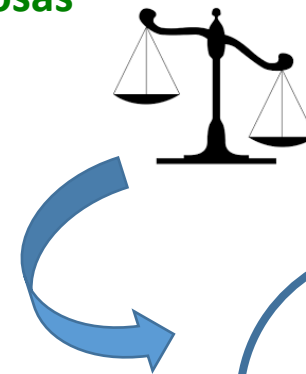
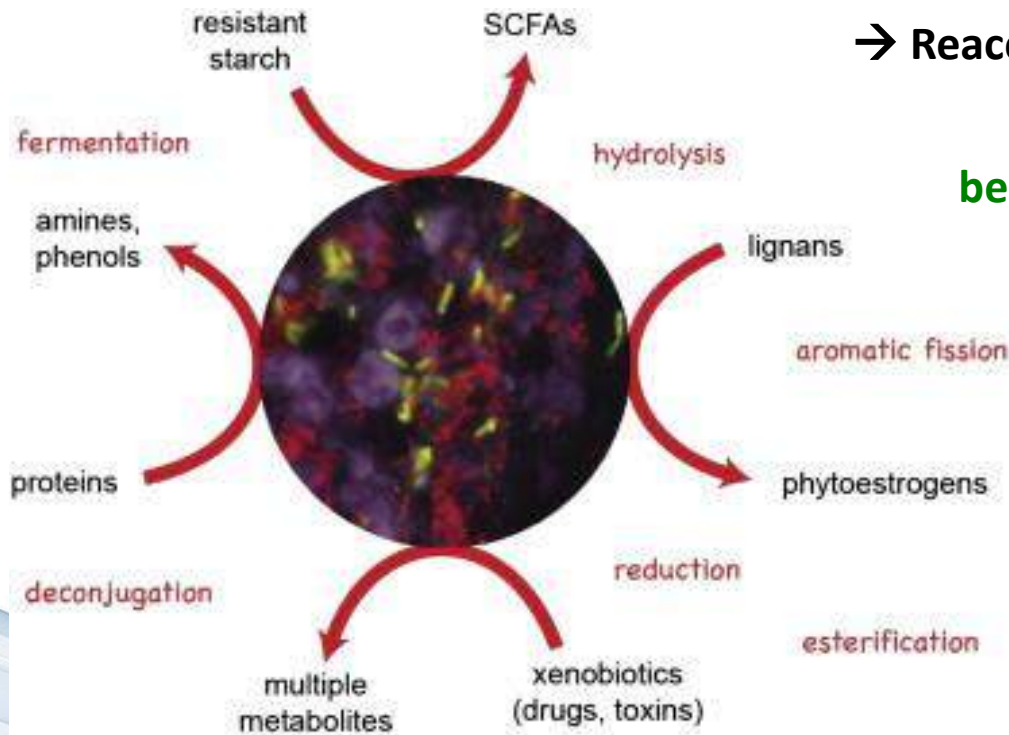


Papel de la microbiota en la salud

→ Reacciones de conversión química

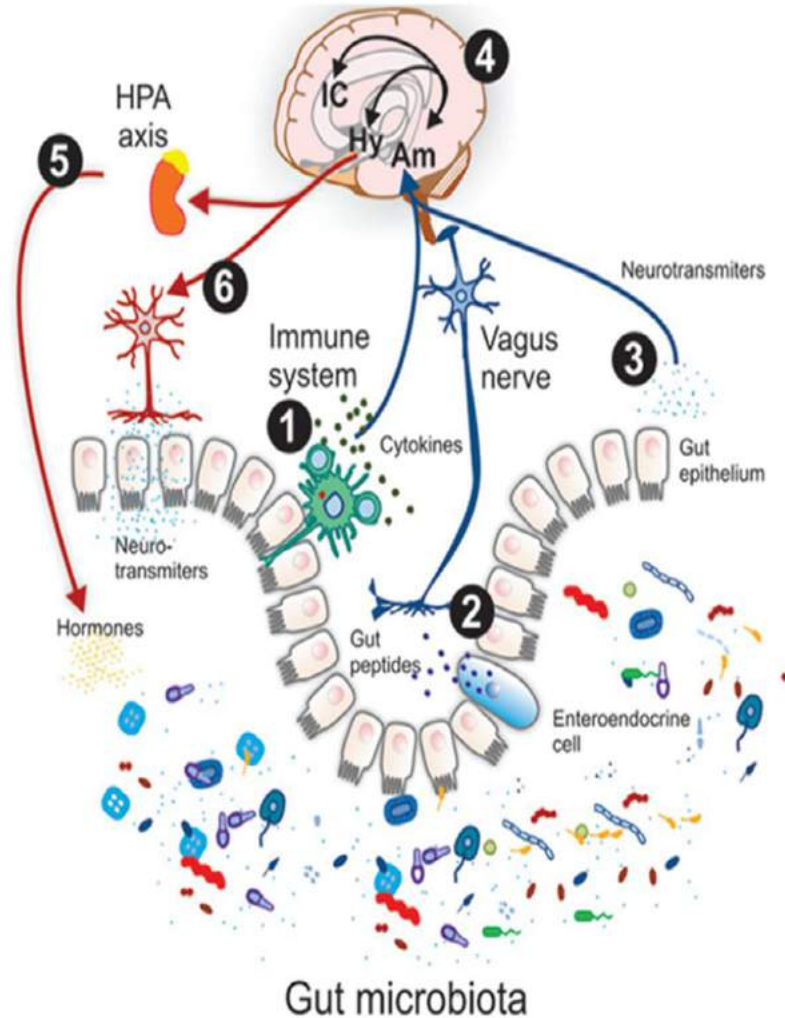
beneficiosas

perjudiciales



Simbiosis
X
Disbiosis

Eje microbiota-intestino-cerebro



- 1 - Linfocitos:** acciones endocrinas o paracrinas
- 2 - Terminales sensoriales neuronales:** activación por péptidos intestinales
- 3 - Neurotransmisores o sus precursores:** efectos endocrinos o paracrinós
- 4 - Activación del hipotálamo, amígdala y cortex insular**
- 5 - Corticosteroides:** liberados como resultado de la activación del eje hipotalámico-pituitario-adrenal (HPA)
- 6 - Activación eferente neuronal:** reflejo colinérgico antiinflamatorio" y/o la activación simpática

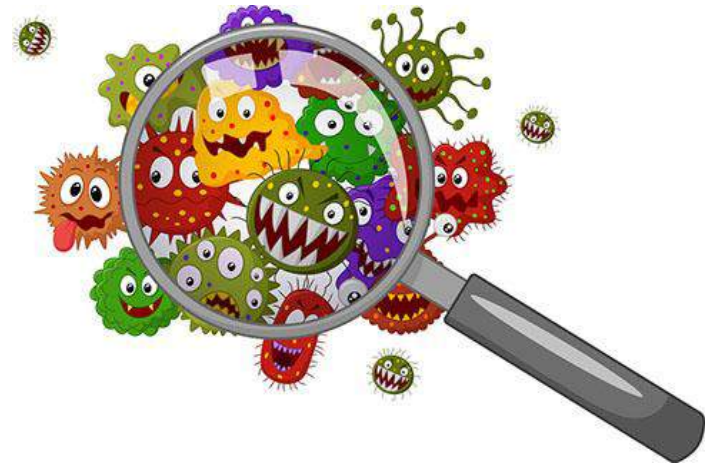
Disbiosis y enfermedades

→ Enfermedades del intestino

- Síndrome de malabsorción
- Enfermedades malignas: cáncer colorrectal
- Enfermedad inflamatoria intestinal
- Síndrome del intestino irritable
- Enfermedades diarreicas
- Infección por Clostridium Difficile

→ Otras enfermedades

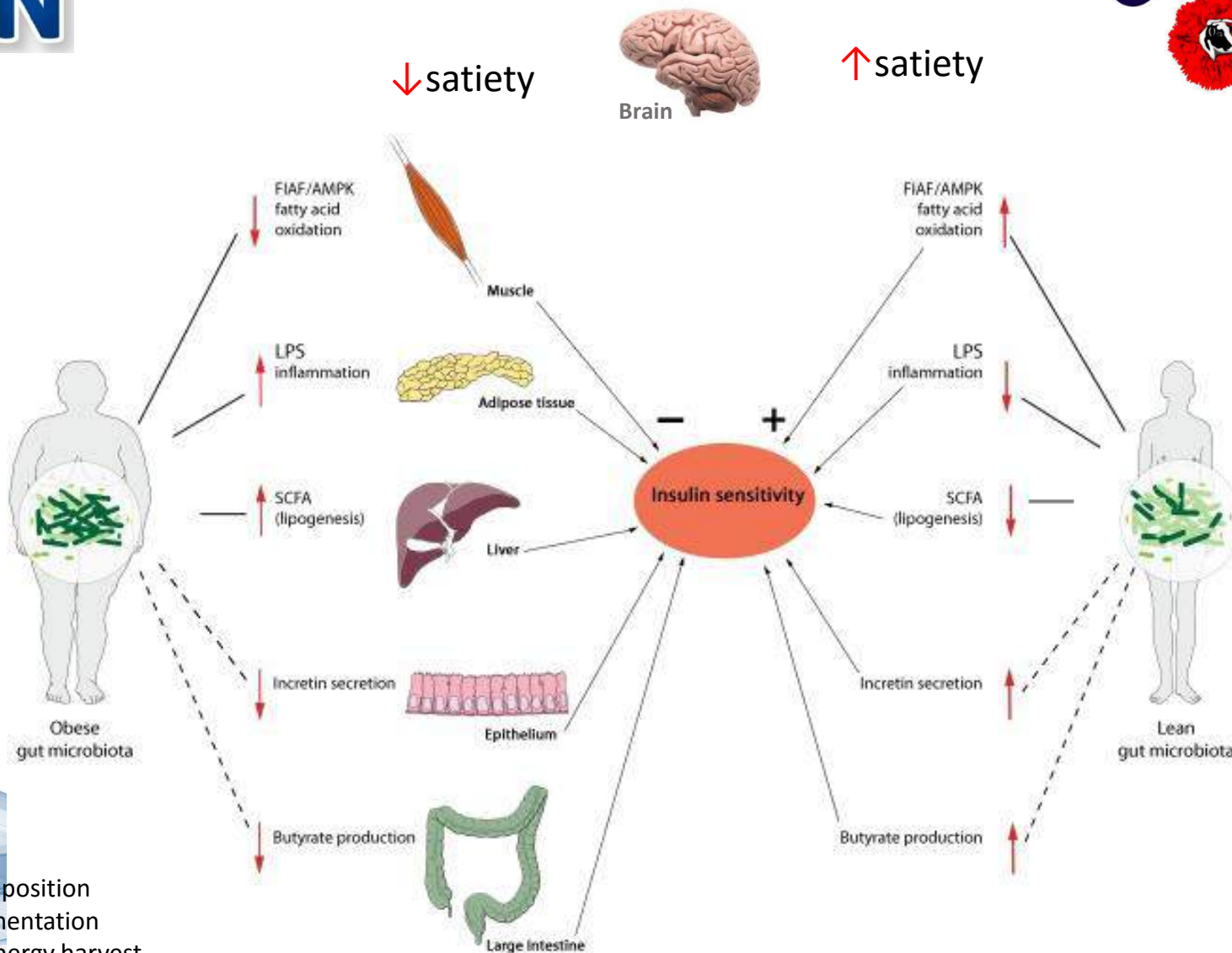
- Malignas: cáncer de hígado, cáncer de mama
- Complicaciones del hígado: cirrosis
- Afecciones alérgicas
- Trastornos autoinmunes (DM1, artritis, etc.)
- Anomalías del eje intestino-cerebro
- Autismo y otros trastornos neurológicos
- **Obesidad y otros transtornos metabólicos**
- Síndrome de fatiga crónica
- Enfermedades periodontales



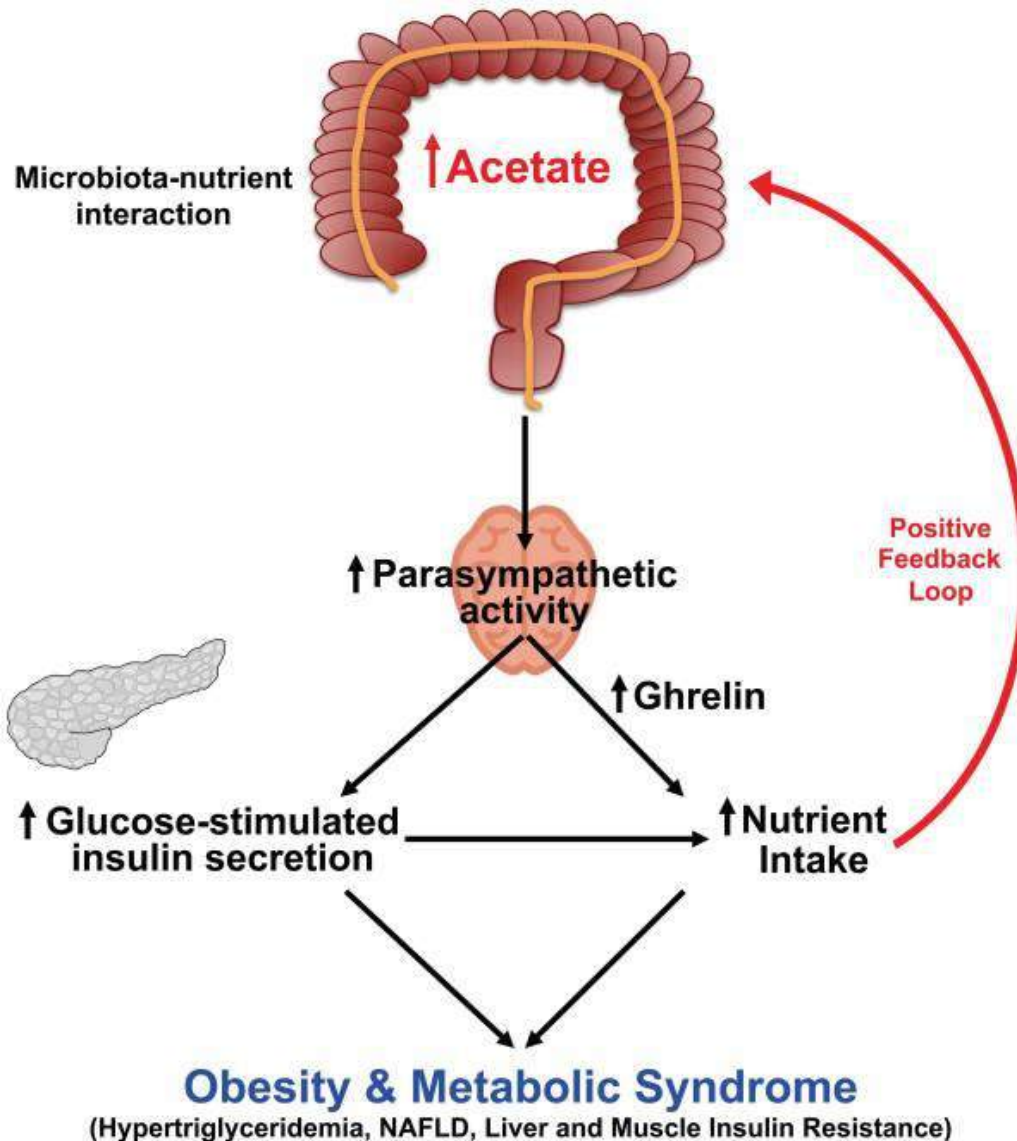
Microbiota intestinal y obesidad



Mayor cantidad relativa de Firmicutes
en relación a los Bacteroidetes



- Altered composition
- Altered fermentation
- Increased energy harvest



El aumento de acetato conduce a un aumento de la ingesta de alimentos, desencadenando un ciclo de **retroalimentación positiva** que impulsa la obesidad y la resistencia a la insulina



Síndrome Metabólico

Tratamiento o prevención de la obesidad

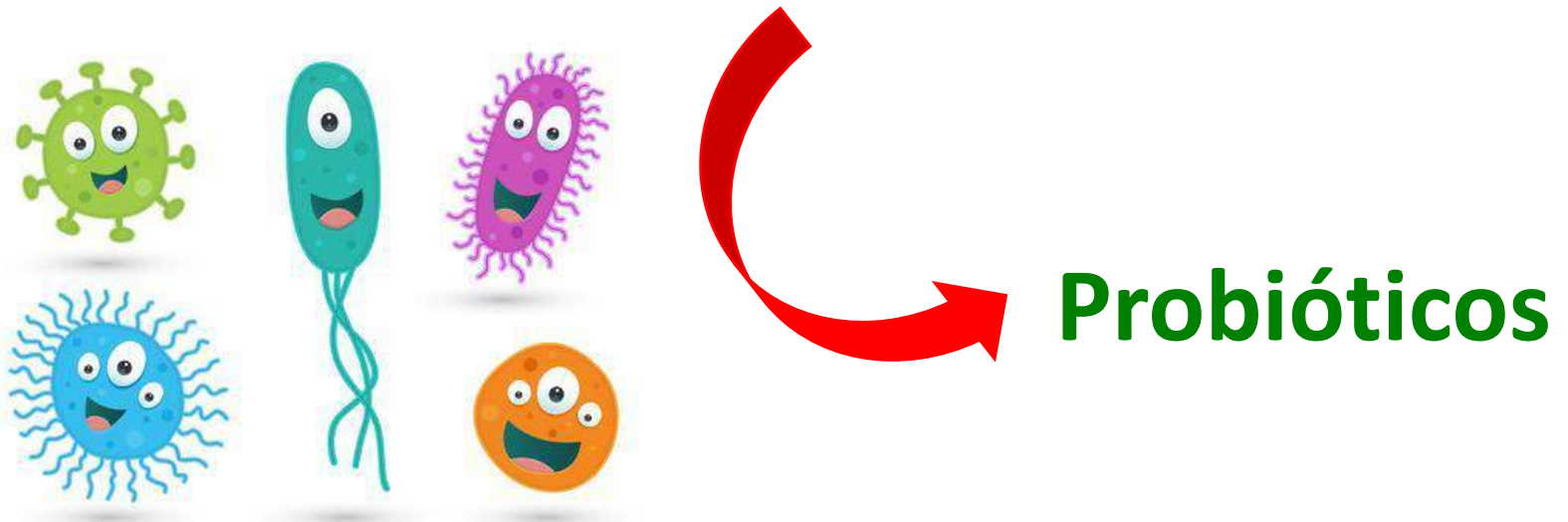


TABLE 1 | Effects of probiotics on gut microbiota of animals and their physiological outcomes.

Probiotics	Animal model	Influence on gut microbiota	Metabolic outcome	Reference
VSL#3	C57J/B67 HFD and ob/ob mice	↓ <i>Firmicutes</i> , ↑ <i>Bacteroidetes</i> , and <i>bifidobacteria</i> . * NC on <i>lactobacilli</i>	↓ Body weight (BW), food intake and adiposity, ↑ insulin sensitivity, glucose tolerance, production of GLP-1 and SCFA butyrate	Yadav et al., 2013
<i>L. curvatus</i> HY7601 and <i>L. Plantarum</i> KY1034	C57BL/6J HFD mice	NC in <i>Firmicutes</i> and <i>Bacteroidetes</i> . Phylum <i>Verrucomicrobia</i> was absent and ↓ <i>Proteobacteria</i> . ↑ Four Species belonging to the <i>Ruminococcaceae</i> and <i>Lachnospiraceae</i> families of the order <i>Clostridiales</i> and phylum <i>Firmicutes</i> . ↑ endogenous <i>B. pseudolongum</i> was found higher	↓ BW and fat accumulation, plasma insulin, leptin, total cholesterol (TC) and liver toxicity biomarkers and adipose tissue, ↓ pro-inflammatory genes in adipose tissue while ↑ fatty acid oxidation genes in liver	Park et al., 2013
<i>L. salivarius</i> UCC118	C57BL/6J HFD mice	NC in <i>Firmicutes</i> , ↑ <i>Bacteroidetes</i> and <i>Proteobacteria</i> and ↓ <i>Actinobacteria</i>	No alteration in metabolic profiles	Murphy et al., 2013
<i>L. salivarius</i> UCC118	C57BL/6J HFD mice	↑ In proportions of <i>Bacteroidetes</i> and <i>Peptococcaceae</i> and significantly ↓ in population of <i>Rikenellaceae</i> and <i>Porphyromonadaceae</i>	↓ BW at appropriate time	Clarke et al., 2013
<i>L. coryniformis</i> CECT5711	C57BL/6J HFD mice	↑ <i>Lactobacillus</i> spp. cluster	↓ Plasma glycaemia, insulin resistance and LPS levels and improves gut permeability.	Toral et al., 2014
<i>L. casei</i> NCDC 19	C57BL/6J HFD mice	NC in total bacterial counts, <i>Eubacterium rectale</i> – <i>Clostridium coccoides</i> and <i>lactobacilli</i> – <i>Enterococcus</i> (LAB) groups. Significant ↑ in population of <i>bifidobacteria</i> was observed	↓ BW, epididymal fat, blood glucose, plasma lipids and leptin; ↑ adiponectin	Rather et al., 2014
<i>L. brevis</i> OK56	C57BL/6J HFD mice	↑ <i>Bifidobacteria</i> population	↓ Body and epididymal fat, ↓ NF-κB activation and LPS production. ↓ Pro-inflammatory cytokines	Kim et al., 2015
<i>L. rhamnosus</i> NCDC17	High fat diet fed and streptozotocin treated Wistar rats	↑ <i>Bifidobacteria</i> and <i>lactobacilli</i> proportions	↑ Glucose tolerance, blood glucose, plasma insulin, glycosylated hemoglobin, FFA, TGs, serum lipids and cholesterol, oxidative stress. ↑ GLP1 and adiponectin, ↓ pro-inflammatory cytokines and propionate in caecum	Singh et al., 2016
Probiotic mixture (<i>L. salivarius</i> 33, <i>L. rhamnosus</i> LMG S-28148, <i>B. animalis</i> subsp. <i>lactis</i> LMG P-28149)	C57BL/6J HFD mice	↑ <i>Bacteroidetes</i> , Restores the adundance of <i>Rikenellaceae</i> ↓ <i>Lactobacillaceae</i> . Probiotic treatment induced a modification in the <i>bifidobacterial</i> population: <i>B. pseudolongum</i> was no longer detected that was present in HFD group. ↓ <i>Lactobacillus</i> – <i>Leuconostoc</i> – <i>Pediococcus</i> group. Restoration of <i>A. muciniphila</i> after treatment	↓ BW and adiposity with improvement in insulin resistance, ↓ adipose tissue inflammation, ↑ dyslipidemia through adipose tissue immune cell-remodeling	Alard et al., 2016
<i>L. plantarum</i> HAC01	C57BL/6J HFD mice	↑ <i>Lachnospiraceae</i> , ↓ <i>Deferribacteraceae</i> , <i>Mucispinitium</i> and <i>Lactobacillaceae</i> with NC in proportion of <i>Firmicutes</i> and <i>Bacteroidetes</i>	↓ BW gain and mesenteric fat weight, blood glucose, TC and triacylglycerol, ↑ lipid oxidative genes	Park et al., 2017

TABLE 2 | Effect of probiotics supplementation on human gut microbiota and their metabolic outcomes.

Probiotic and subject	Study Design	Influence on gut microbiota	outcome	Reference
<i>L. rhamnosus</i> GG, ATCC 53103 pregnant women and children	Double-blind placebo controlled Mothers: 4 weeks before deliver Children: 6 months after birth	No clear cut study on gut/fecal microbiota	The authors proposed that the reduction in BW was due to positive modulation of gut microbiota by probiotic during the critical development period.	Luoto et al., 2010
<i>L. salivarius</i> Ls-33 obese adolescents	Double-blind placebo controlled 12 weeks	↑ <i>Bacteroides-Prevotella-Parapharymonas</i> group to <i>Firmicutes</i> belonging bacteria, including <i>Clostridium</i> cluster XIV, <i>Blautia coccoides</i> , <i>Eubacteria rectale</i> and <i>Roseburia intestinalis</i> group. NC of ingestion on <i>Clostridium</i> cluster IV, <i>F. prausnitzii</i> , <i>Enterobacteriaceae</i> , <i>Enterococcus</i> , <i>Lactobacillus</i> , and <i>Bifidobacterium</i> group	Feeding strain might have modified the fecal microbiota in obese adolescents by a mechanism that is not associated to metabolic syndrome.	Larsen et al., 2013
Probiotic (<i>S. thermophilus</i> , <i>L. plantarum</i> , <i>L. acidophilus</i> , <i>L. rhamnosus</i> , <i>B. lactis</i> , <i>B. longum</i> and <i>B. breve</i>) and <i>Bofutsushosan</i> herb, subjects having BMI > 25 kg/m ² and waist circumference > 85 cm were only included in the study	Double-blind, placebo controlled 8 weeks	↑ <i>B. breve</i> , <i>B. lactis</i> , <i>L. rhamnosus</i> , and <i>L. plantarum</i> . ↓ <i>Firmicutes/Bacteroidetes</i> ratio was also observed upon treatment.	↓ Weight, waist circumference and ↑ in HDL-cholesterol. Change in body composition is positively related to levels of LPS and <i>L. plantarum</i>	Lee et al., 2014
<i>L. salivarius</i> UBL S22 and Prebiotic Fructooligosaccharide healthy human volunteers	Double-blind, placebo controlled 6 weeks	↑ <i>Lactobacilli</i> and ↓ <i>E. coli</i> after intervention period.	Improvement in serum TGs and lipid profile with ↓ in inflammatory cytokines	Rajkumar et al., 2015

ORIGINAL ARTICLE

Regulation of abdominal adiposity by probiotics (*Lactobacillus gasseri* SBT2055) in adults with obese tendencies in a randomized controlled trial

Y Kadooka¹, M Sato², K Imaizumi², A Ogawa¹, K Ikuyama³, Y Akai³, M Okano³, M Kagoshima⁴ and T Tsuchida⁵



- ensayo de intervención doble ciego, aleatorizado, controlado con placebo
- sobrepeso y obesos
- 33 a 63 años de edad
- grupo control (n=44) y grupo placebo (n=44)
- *Lactobacillus gasseri* SBT2055 en leche fermentada
- 12 semanas

ORIGINAL ARTICLE

Regulation of abdominal adiposity by probiotics (*Lactobacillus gasseri* SBT2055) in adults with obese tendencies in a randomized controlled trial

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Table 2 Abdominal fat areas

Parameters	Values	Groups	W0	W12		Group-by-time Group
Visceral fat (cm ²)	Measured	Active	127.3 (119.7, 134.9)	121.5 (130.2, 112.8)	**	##
		Control	119.3 (112.8, 125.8)	120.7 (114.0, 127.4)		
	Change from W0	Active	—	−5.8 (−10.0, −1.7)		##
		Control	—	1.4 (−2.2, 5.1)		
Subcutaneous fat (cm ²)	Measured	Active	222.2 (196.9, 247.5)	214.8 (189.4, 240.3)	**	#
		Control	227.8 (204.4, 251.2)	226.5 (204.0, 249.1)		
	Change from W0	Active	—	−7.4 (−11.6, −3.1)		#
		Control	—	−1.3 (−4.9, 2.4)		
Total fat (cm ²)	Measured	Active	349.5 (322.5, 376.6)	336.3 (309.5, 363.2)	**	##
		Control	347.1 (321.3, 372.9)	347.2 (321.9, 372.6)		
	Change from W0	Active	—	−13.2 (−20.5, −6.0)		##
		Control	—	0.1 (−5.7, 6.1)		

Values are means (95% confidence interval (CI)). Within-group comparisons between week 0 (W0) and W12: ** $P < 0.01$. Between-group comparisons: # $P < 0.05$, ## $P < 0.01$. Group-by-time interaction ('Group-by-time'): # $P < 0.05$, ## $P < 0.01$. Group-effect ('Group'): not significant.

The Additional Effects of a Probiotic Mix on Abdominal Adiposity and Antioxidant Status: A Double-Blind, Randomized Trial

Aline Corado Gomes¹, Rávila Graziany Machado de Sousa¹, Patrícia Borges Botelho¹, Tatyane Letícia Nogueira Gomes¹, Patrícia Oliveira Prada², and João Felipe Mota¹

- **Ensayo aleatorizado, doble ciego**



- **sobrepeso y obesas**
- **20 a 59 años de edad**
- **grupo control (n=21) y grupo placebo (n=22)**
- **8 semanas**
- **mix de probióticos: Lactobacillus acidophilus LA-14, Lactobacillus casei LC-11, Lactococcus lactis LL-23, Bifidobacterium bifidum BB-06, y Bifidobacterium lactis**

TABLE 1 Body composition before and after treatment with dietary intervention or a dietary intervention plus probiotic mix in women with overweight and obesity

	Dietary intervention (n = 22)				Dietary intervention + probiotic mix (n = 21)					Intervention effect	
	Baseline	Week 8	Mean change	P ^a	Baseline	Week 8	Mean change	P ^a	P ^b	Effect (95% CI)	P ^c
BMI (kg/m ²)	33.34 ± 4.69	32.61 ± 4.53	-0.72	0.00	31.70 ± 3.90	31.24 ± 3.96	-0.45	0.01	0.41	-0.49 (-3.24 to 2.25)	0.71
Weight (kg)	83.55 ± 13.44	82.59 ± 13.00	-0.95	0.02	76.58 ± 9.94	75.60 ± 10.24	-0.98	0.02	0.18	-3.64 (-11.24 to 3.95)	0.33
WC (cm)	97.50 ± 10.40	94.18 ± 10.24	-3.32	0.00	93.95 ± 8.37	88.80 ± 7.26	-5.14	0.00	0.33	1.81 (0.13 to 3.50)	0.03
Body fat (%)	48.82 ± 4.39	48.03 ± 4.18	-0.79	0.02	48.56 ± 4.28	47.37 ± 5.00	-1.19	0.00	0.91	-1.09 (-3.91 to 1.72)	0.43
FM (kg)	39.73 ± 8.23	39.05 ± 7.35	-0.68	0.29	36.30 ± 7.32	34.96 ± 7.52	-1.33	0.00	0.60	-1.93 (-7.02 to 3.15)	0.44
FFM (%)	49.68 ± 4.33	50.36 ± 4.09	0.68	0.04	49.79 ± 4.09	50.92 ± 4.76	1.13	0.00	0.79	-0.94 (-1.79 to 3.69)	0.48
FFM (kg)	41.37 ± 6.74	41.46 ± 6.53	0.09	0.77	37.86 ± 3.65	38.20 ± 4.20	0.33	0.26	0.00	-1.13 (-4.68 to 2.41)	0.52
Android (%)	54.36 ± 3.35	53.05 ± 3.23	-1.31	0.00	54.80 ± 3.36	53.50 ± 4.14	-1.30	0.02	0.98	0.03 (-2.32 to 2.39)	0.97
Gynoid (%)	53.98 ± 5.31	52.75 ± 5.42	-1.22	0.00	53.95 ± 4.66	53.13 ± 4.70	-0.81	0.00	0.56	-1.92 (5.12 to 1.26)	0.23
WHR	0.61 ± 0.06	0.59 ± 0.06	-0.02	0.00	0.60 ± 0.06	0.57 ± 0.05	-0.03	0.00	0.87	0.01 (0.00 to 0.02)	0.02
Conicity index	1.23 ± 0.07	1.19 ± 0.07	-0.03	0.00	1.22 ± 0.07	1.16 ± 0.06	-0.05	0.92	0.91	0.02 (0.00,0.04)	0.03

Values are presented as mean ± standard deviation.

^aDifference between baseline and end point. *P* value obtained from paired *t*-test/Wilcoxon matched-pairs signed-rank test for the within-group comparisons.

^bHomoscedasticity test between groups on baseline.

^cObtained from unpaired Student's *t*-test or Mann-Whitney test, as statistical difference between changes (DI vs. DI + P). The effect of the probiotic mix associated with dietary intervention on the variables measured was examined using ANCOVA with adjustment for the screening/baseline observation.

FFM, free-fat mass; FM, fat mass; WC, waist circumference; WHR, waist-to-height ratio.

↓ de la adiposidad abdominal

TABLE 5 Oxidative and inflammatory parameters before and after treatment with dietary intervention or a dietary intervention plus probiotic mix in women with overweight and obesity

	Dietary intervention (n = 22)				Dietary intervention + probiotic mix (n = 21)					Intervention effect	
	Baseline	Week 8	Mean change	P ^a	Baseline	Week 8	Mean change	P ^a	P ^b	Effect (95% CI)	P ^c
CAT (U/mg)	8.10 ± 2.01	7.17 ± 2.05	-0.92	0.15	8.20 ± 1.26	7.71 ± 1.15	-0.49	0.65	0.06	0.30 (-0.50 to 0.11)	0.44
SOD (U/mg)	5.88 ± 1.94	6.28 ± 1.90	0.40	0.55	5.35 ± 2.50	7.31 ± 6.29	1.95	0.00	0.10	-0.62 (-2.79 to 1.54)	0.56
GPx (U/mg)	0.42 ± 0.12	0.34 ± 0.10	-0.07	0.00	0.32 ± 0.13	0.37 ± 0.11	0.05	0.04	0.31	-0.12 (-0.22 to 0.02)	0.01
MDA (nmol/mL)	0.16 ± 0.04	0.16 ± 0.04	0.00	0.83	0.18 ± 0.25	0.21 ± 0.28	0.02	0.39	0.00	-0.01 (-0.13 to 0.10)	0.81
LPS (EU/mL)	0.85 ± 0.44	0.95 ± 0.60	0.09	0.49	0.67 ± 0.11	0.65 ± 0.11	0.02	0.34	0.00	0.08 (-0.26 to 0.44)	0.62
IL-10 (pg/mL)	4.65 ± 4.34	4.53 ± 3.14	-0.12	0.88	3.57 ± 1.66	3.96 ± 1.84	0.39	0.15	0.00	-1.29 (-3.34 to 0.74)	0.20
IL-6 (pg/mL)	1.35 ± 1.81	0.80 ± 1.81	-0.54	0.07	0.58 ± 0.27	0.58 ± 0.31	0.00	0.94	0.00	-1.98 (-4.15 to 0.18)	0.07
TNF-α (pg/mL)	2.49 ± 0.89	2.07 ± 0.78	-0.42	0.07	1.72 ± 0.44	1.88 ± 0.37	0.16	0.24	0.00	-0.84 (-1.47 to -0.20)	0.01
ADIPO (ug/mL)	35.59 ± 20.76	38.44 ± 29.93	2.84	0.65	33.72 ± 29.78	36.33 ± 27.46	2.61	0.63	0.11	3.92 (-15.27 to 23.11)	0.68

Values are presented as mean ± standard deviation.

^aDifference between baseline and end point. *P* value obtained from paired *t*-test/Wilcoxon matched-pairs signed-rank test for the within-group comparisons.

^bHomoscedasticity test between groups on baseline.

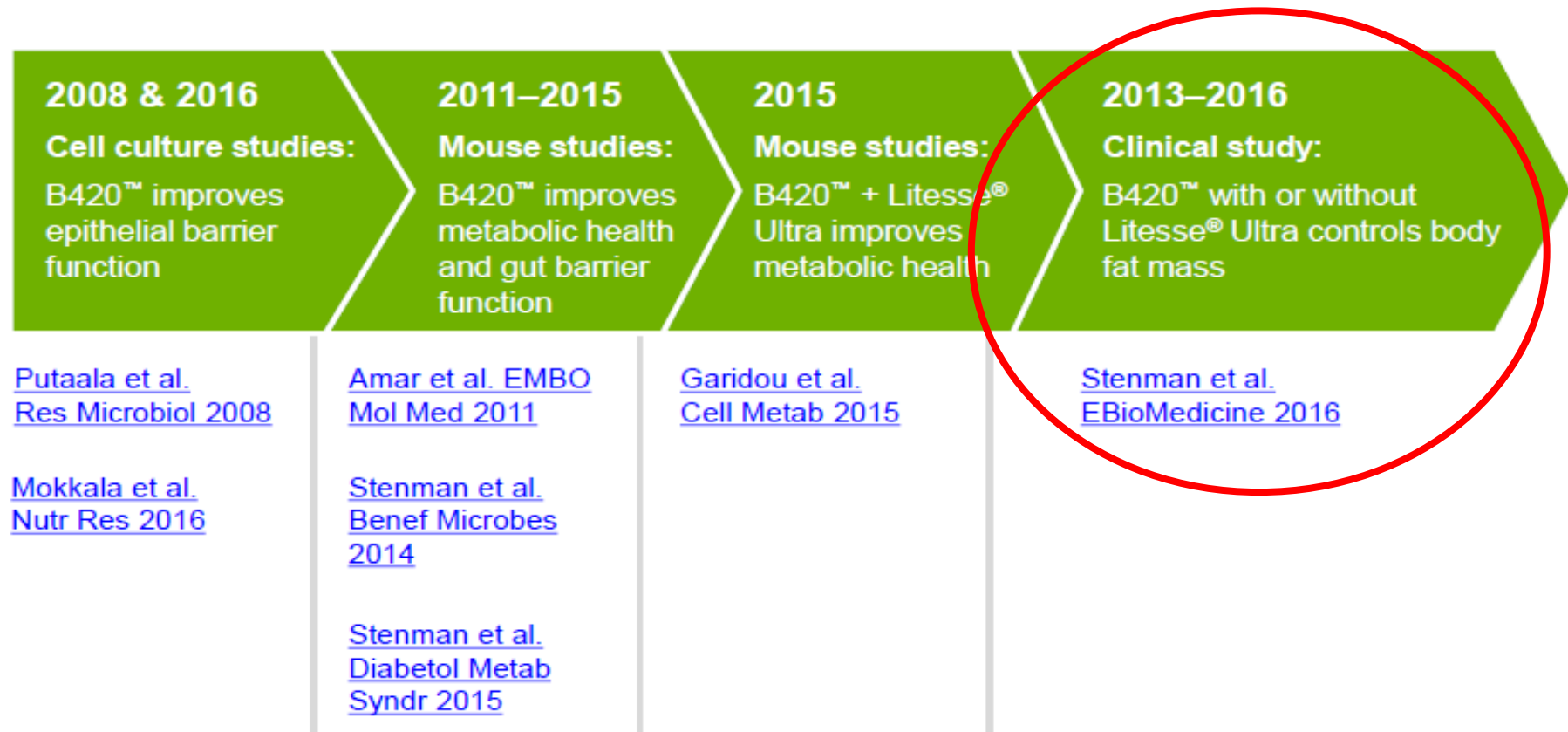
^cObtained from unpaired Student's *t*-test or Mann-Whitney test, as statistical difference between changes (DI vs. DI + P). The effect of the probiotic mix associated with dietary intervention on the variables measured was examined using ANCOVA with adjustment for the screening/baseline observation.

ADIPO, adiponectin; CAT, catalase; SOD, superoxide dismutase; GPx, glutathione peroxidase; IL, interleukin; LPS, lipopolysaccharide; MDA, malondialdehyde; TNF-α, tumor necrosis factor-α.

↑ de la actividad enzimática antioxidante

Scientific Documentation

Bifidobacterium animalis ssp. *lactis* 420 (B420™) for metabolic health – a decade of research





- ensayo de intervención doble ciego, aleatorizado, controlado con placebo
- 255 participantes
- sobrepeso y obesos
- 18 a 65 años de edad

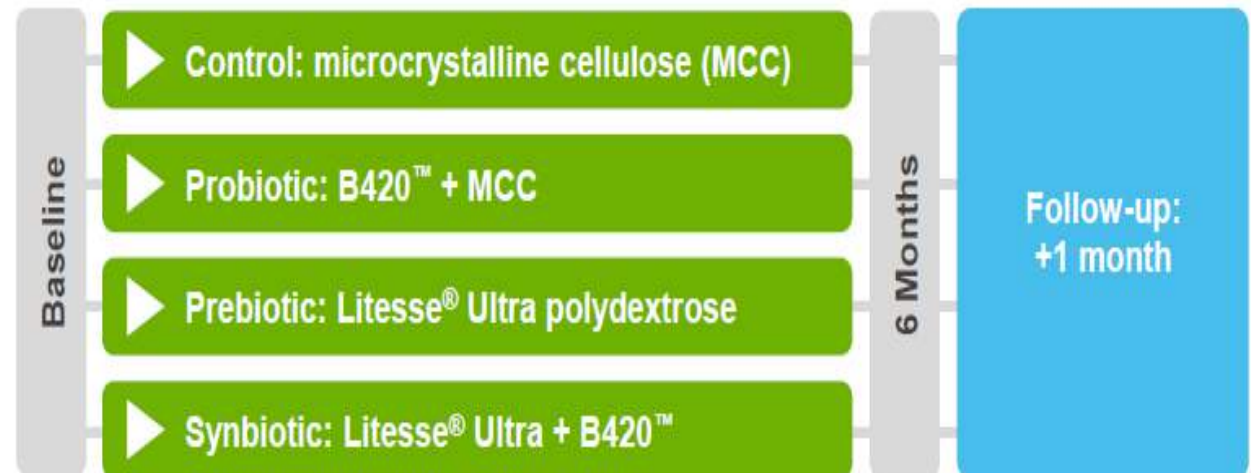


Research Paper

Probiotic With or Without Fiber Controls Body Fat Mass, Associated With Serum Zonulin, in Overweight and Obese Adults—Randomized Controlled Trial☆☆☆



Lotta K. Stenman ^{a,*}, Markus J. Lehtinen ^a, Nils Meland ^b, Jeffrey E. Christensen ^c, Nicolas Yeung ^a, Markku T. Saarinen ^a, Michael Courtney ^c, Rémy Burcelin ^c, Marja-Leena Lähdeaho ^d, Jüri Linros ^e, Dan Apter ^f, Mika Scheinin ^g, Hilde Kloster Smerud ^b, Aila Rissanen ^h, Sampo Lahtinen ^a



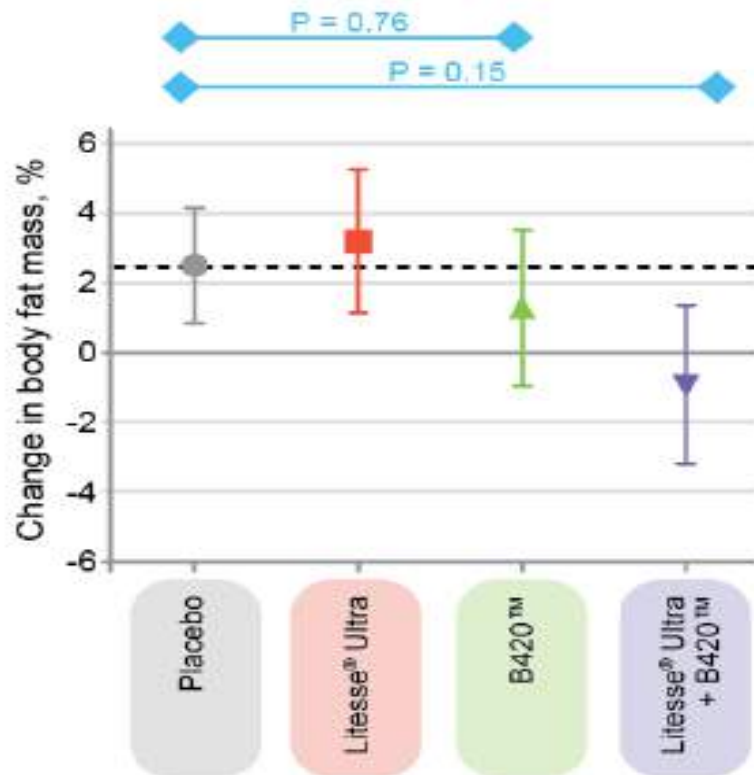
- **B420™**
 - 12 g microcrystalline cellulose
 - 10^{10} CFU/day *B. lactis* 420
- **Litesse® Ultra**
 - 12 g Litesse® Ultra polydextrose
- **Litesse® Ultra + B420™**
 - 12 g Litesse® Ultra polydextrose
 - 10^{10} CFU/day *B. lactis* 420
- **Placebo**
 - 12 g microcrystalline cellulose (no energy; less fermented)



Smoothie contains 130 kcal/day

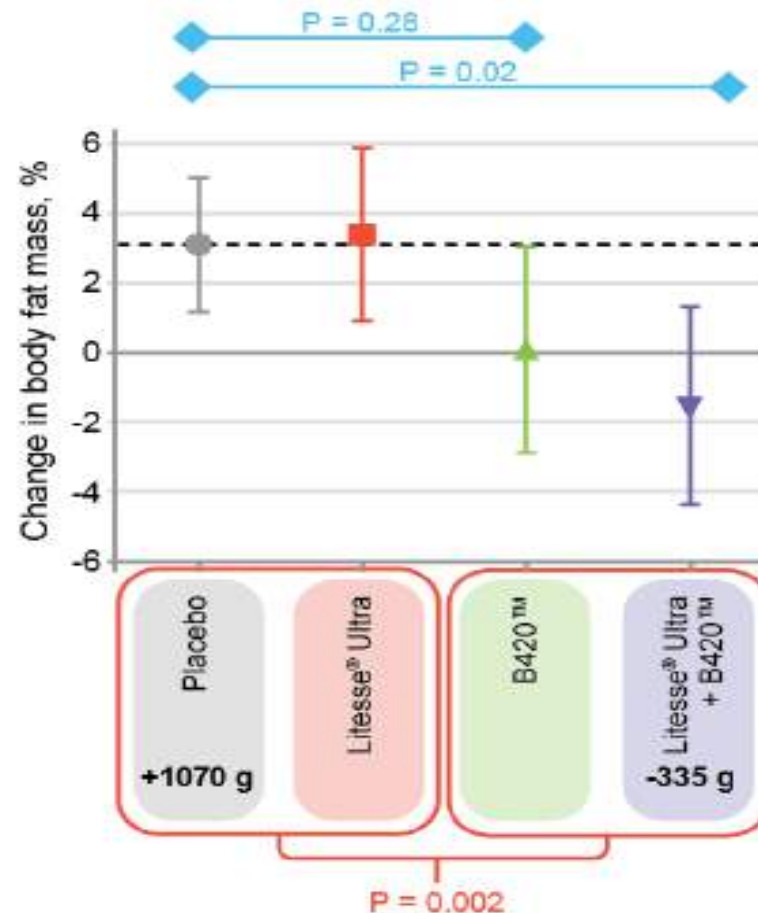
Intention to Treat

Overall ANCOVA $P=0.46$



Per Protocol

Overall ANCOVA $P=0.095$

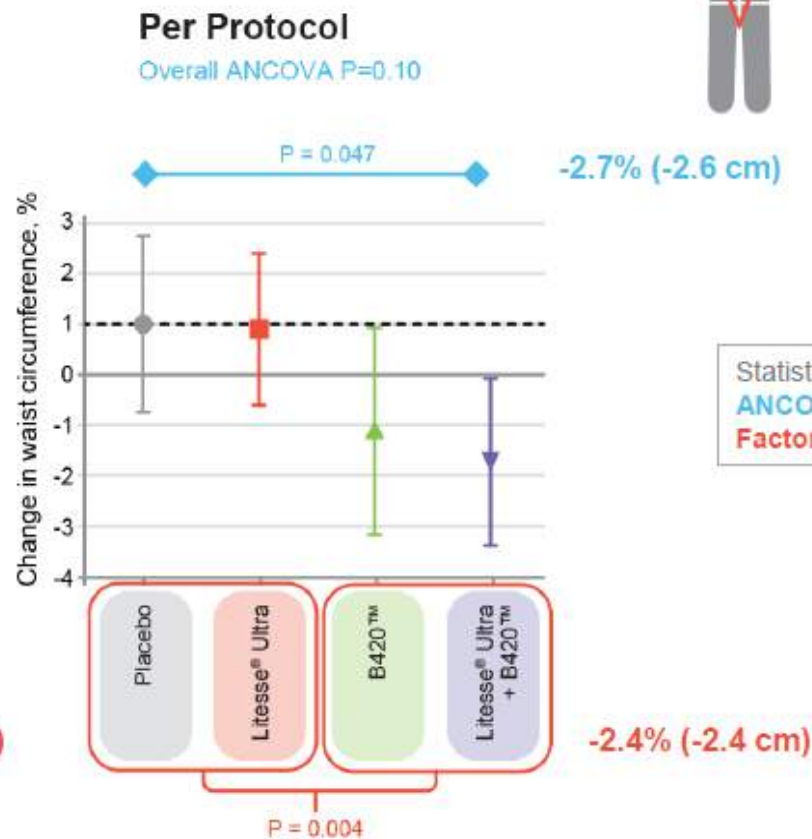
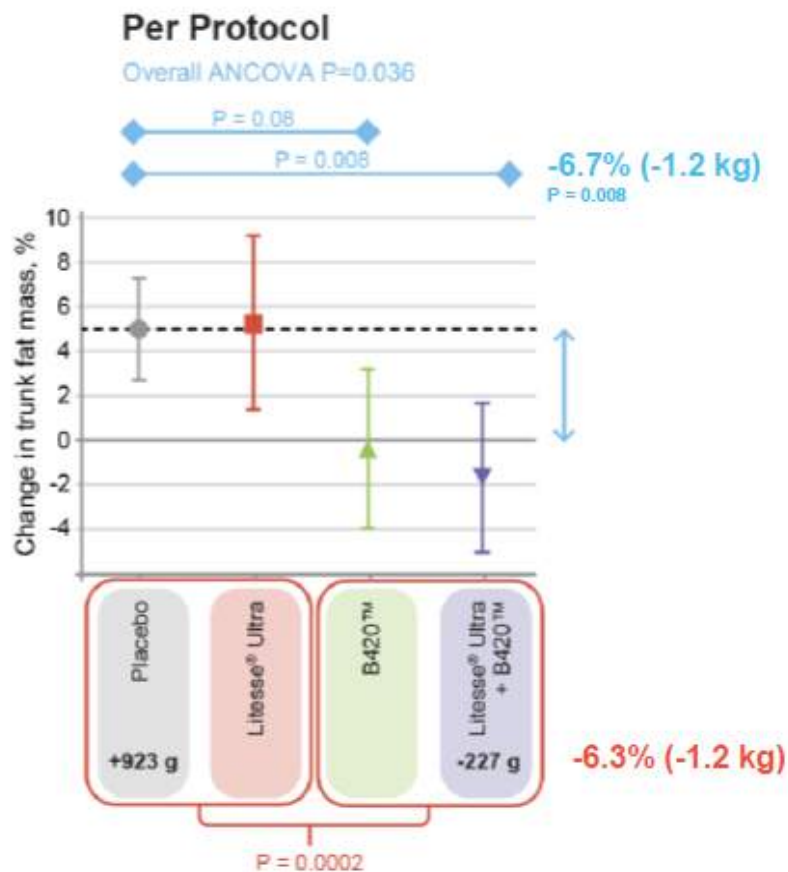


Stenman et al. EBioMedicine 2016

→ Reducción en 4,5% (-1,4 Kg) de grasa corporal comparado con placebo

→ Efecto concentrado en el área abdominal

Effect concentrated to abdominal area

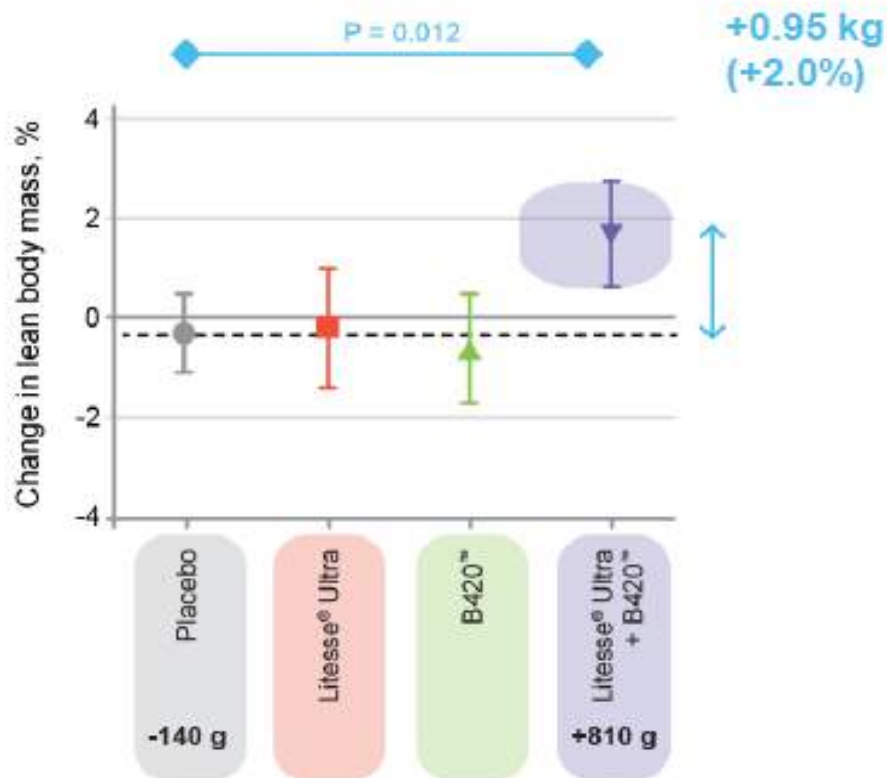


Statistical analyses:
ANCOVA values
Factorial analysis

Lean body mass

Per Protocol population

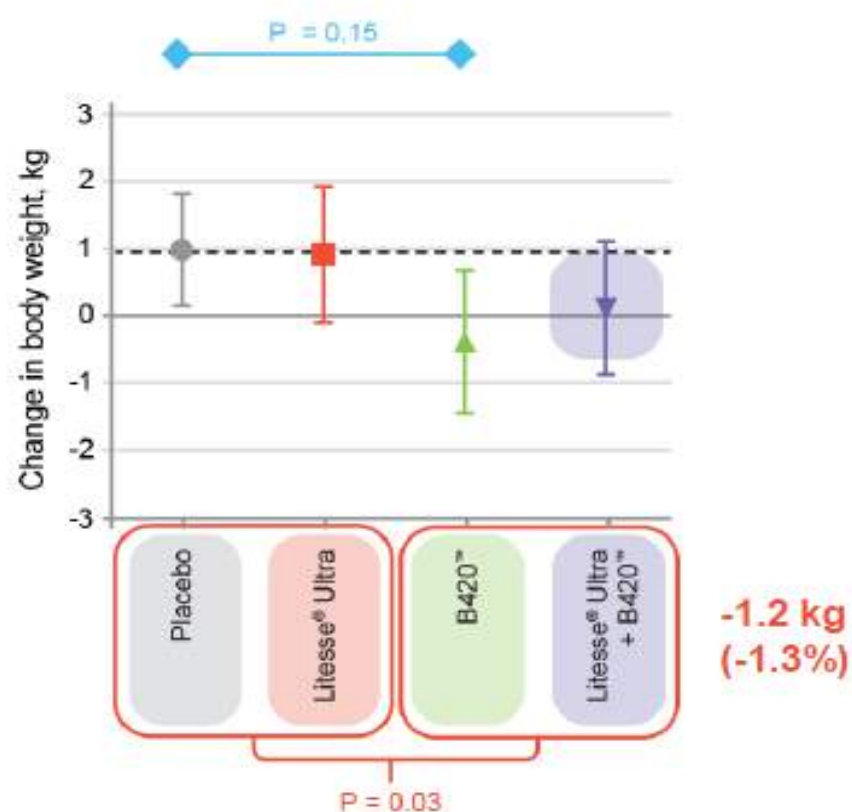
Overall ANCOVA $P=0.30$



Body weight

Per Protocol population

Overall ANCOVA $P=0.13$

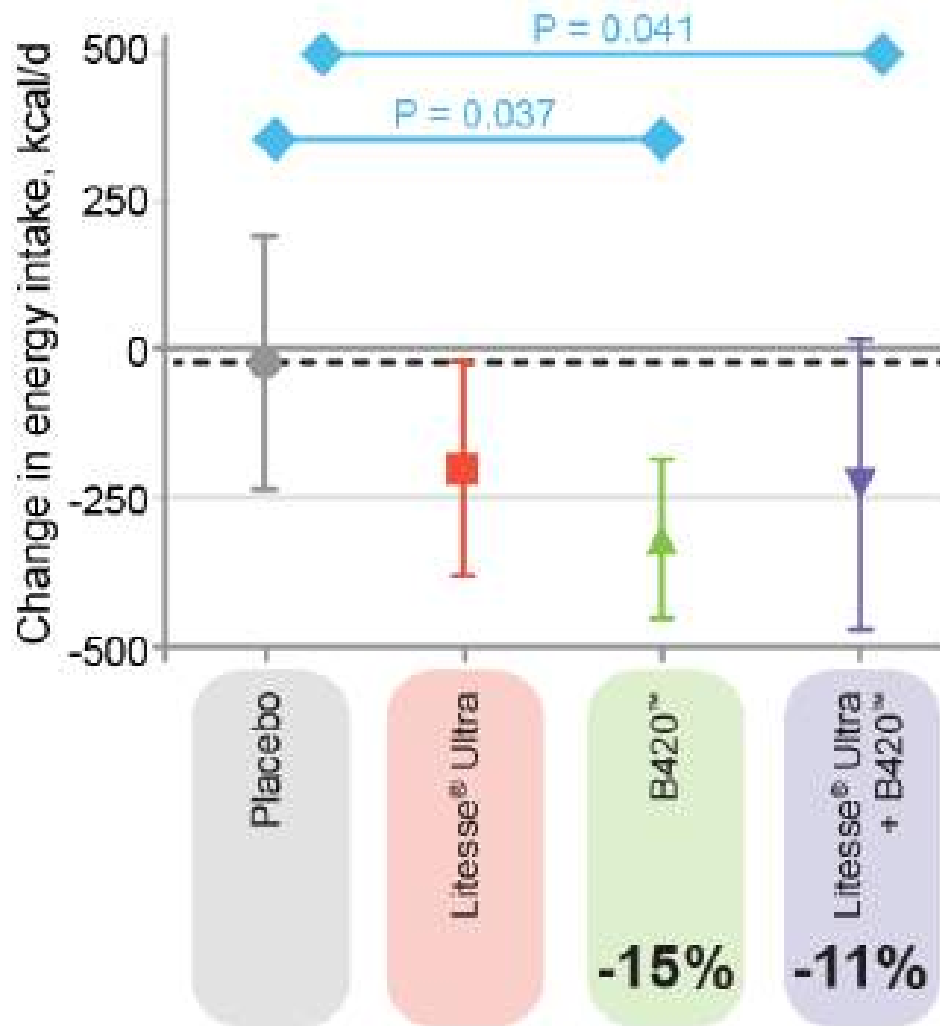


Stenman et al. EBioMedicine 2016

→ Reducción en 1,2 Kg de peso corporal comparado con placebo

Energy intake

Overall ANCOVA $P=0.005$



→ B420™ solo o con polidextrosis reduce la ingesta de energia

Data from 5-day food diaries, Per Protocol population

Probióticos



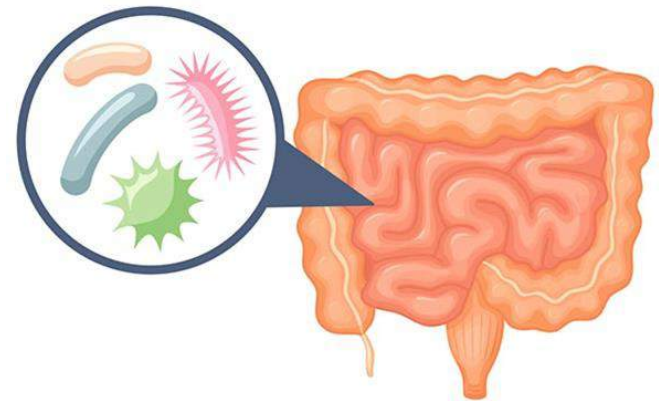
**Tratamiento o prevención
de la obesidad**



Gracias por su atención!



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INTERACCIÓN DE LOS PROBIÓTICOS CON LA MICROBIOTA INTESTINAL Y OBESIDAD



Dra. Bárbara Emo Peters