

(1–5 Mb ecDNA al^l kn⁺ n a⁺ d⁺ ble min⁺ e⁺) (Pallen et al. 2018; Shiba et al. 2012; Tanne et al. 2017). Circular DNA can de⁺ ef⁺ m⁺ i e⁺ i h l l e⁺ n⁺ e⁺ ence h⁺ m⁺ l⁺ g⁺, h⁺ e⁺ e⁺, highl⁺ e⁺ e⁺ i i e⁺ gen⁺ mic⁺ egi⁺ n⁺, . . . ch a⁺ he⁺

cicla DNA, ecie d' n' f' ma a high a e, n' c' n ain
aci e e lica i' n' igin and, he ef' e, he c' n mbe' f
he e, ecie in m' he cell ill emain a ic' dec ea e
i h age. A c' i' l' l' c' me' f' he e dr' e ence in f' ma-
i' n' eed, e lica i' n' ca aci and a mme ic' e en i' n
i' ha he di e, i' f ci c la DNA, ecie b' e ed in
l' ng cell i' high, b' he c' n mbe' f each indi id al
ci cle and, he ef' e, he hen' ic im ac i' l'. A cell
age, 'n' l' a' b' e' f ci c la DNA can acc' m' la e b' ha

megaba e ff DNA, gene- ich ch 'm ! 'me c 'n ib . e m 'e . . he . . al le el ff ci c la DNA in heal h h man i . . e (M ille e al. 2018). F he m 'e, *TTN*(i in), he m ! . an c ibed lein-c ding gene in m . cle i . . e, i al . he la ge . . d ce ff ci c la DNA e gene (M ille e al. 2018). F ! mali ing hi idea, e . . gge . ha gene hich ha e e ! ed . be ind ced in e . . n e . . a ic . -la en i . nmen al c 'ndi i 'n a e e cellen candida e f ! ada . i e am li ca i 'n, and ha . im 1 c 'nnec ing ci c -la DNA f ! ma i 'n . an c i . i al ind c i 'n i a cle e mean b . hich cell c ! ld gain he ma im m chance ff acc. m la ing . . ef. 1 ci c la DNA, a he . han . nhel f 1 ! nega i e . ecie .

B i . elf, an ada i e hen ! e in an indi id al aged cell i . f li le . . e if he ca . al ci c la DNA i . el . hl e ained in he m ! he cell, a . . l d be he ca e if a . mme ic . eg ega i 'n i main ained, and e m . . c 'n ide h ! ci c la DNA acc. m la i 'n i . an la ed in 'a he i -able ad an age. Fi . . , nce ci c la DNA ha acc. m la ed, eg ega i 'n can be ela ed . nde . . e . all ! ing ci cle i h e lica i 'n ! igin . . ! aga e a high c ! . n mbe in he . . la i 'n (Fig. 1, . . 5a). Thi . elea e ff he a . mme ic . eg ega i 'n . . em . nde hea . . e . ha been b e ed and e . e en . a gene al e . . n e . . ignalling f ! m he cell . all in eg i . . a h a (Baldi e al. 2017). Sec . . ndl , acc. m la i 'n f high le el ff ci c la DNA inc ea e . he chance ff ch 'm ! 'me e-in eg a i 'n and, he ef ! e, e . . a i 'n f n ! mal he i abili f ! . he am li -ed allele (Fig. 1, 5b). S ch ada i e ch 'm ! 'mal e -in eg a i 'n e en . ha e been e ea edl b e ed, al h ! gh i i . nclea . he he . he ha ened in aged cell (Be e le e al. 1984; Be e e al. 2015; Demeke e al. 2015; D kin e al. 2012; Gale e e al. 2011; K che e al. 2020; La e e al. 2018; V g e al. 2004).

The idea ha a . b- . . la i 'n . ade . h ! . - e m g ! . h f ! ada i e ca aci i f ! mali ed in be -hedging (c 'nci el e ie ed in (Le e al. 2012)), and face . . f ageing ha . . i h a be -hedging m 'del ha e been dem . . a ed e . e i-

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