So, good evening everyone and thank you organizers for giving me this opportunity. So, today I am going to discuss on unleashing the potential of probiotic fissile

bacterium

prosonids to optimize the PD-1 inhibitor efficacy in non-small cell lung cancer treatment.

Coming to the background, if you see in the locally advanced non-small cell lung carcinoma,

the survival rate has been ranged from 13 to 36 percent.

On average for stage 3 non-small cell lung cancer, the 5 year overall survival rate is 30 percent.

We have made a lot of improvements in the treatments and we have incorporated immunotherapy also

in the locally advanced lung carcinoma management.

And in one such Chinese study that is Neo-Tarch, what they have noted was by incorporating immunotherapy

in locally advanced non-small cell lung cancer.

They could achieve the major pathological response of up to 50 percent.

What do you, I mean by major pathological response was, pathological response rate in

more than 90 percent of the cells is called as major pathological response and such rate

was observed in almost 50 percent of the patient.

And there is an even free survival benefit also by adding this immunotherapy in the perioperative setting.

So, again there are multiple studies which have shown that there is an association between

the immunotherapy and the gut microbiota.

There are patients who are responders to the immunotherapy based on what gut microbiota

they have and there are certain patients who are classified as non-responders based on

what gut microbiota they have.

And similarly, depending on the presence of this gut microbiota, there is an altered

incidence of colitis because of the immunotherapy.

So, one such bacterium, today I wanted to discuss is physically bacterium prosonates.

So, first I just wanted to highlight what is the metabolism of like how this probiotic

work and how it will help in the immune system.

So, if you look at this right side walla, the KY and A and 113 A, these are the tryptophan

metabolites.

And when this tryptophan metabolites bind to the AHR complex, it will modify the tumor

micro-environment, it will promote the tumor growth and it will suppress the CD8 T cell

activation.

These are the two points you should remember.

So, when the tryptophan metabolites bind to the AHR complex, it promotes the tumor growth

and it inhibits the CD8 T cell.

So, what we wanted to achieve was to remove this tryptophan metabolites from the tumor.

So, how we can achieve of removing this tryptophan metabolites are the metabolite component of

this facility bacterium prosonates is valeric acid.

And this valeric acid in turn up regulates the run X3 gene and it will in turn causes

the in decreased expression of the interleukin IV induced gene 1, which is the major step

in the catabolism of the tryptophan.

So, with the help of this valeric acid, we can decrease the level of interleukin IV induced gene 1, which in turn decrease or down regulate the catabolism of the tryptophan.

Thereby, there is a decreased availability of the tryptophan metabolites, which in turn

causes the activation of the CD8 T cell.

So, this is the basic metabolic effect of how this facility bacterium prosonates will

help in increasing the efficacy of the PDL1 inhibitors.

So, coming to the methods and methodology, patient has been received knee adjuvant immunotherapy

and chemotherapy and they have classified as responders and non-responders depending

on the presence of major pathological response present or not.

And from these patients, either both from the responders and non-responders, they have

collected the tumor and feces samples and they have underwent the metagenomic sequencing

to study this microbiota.

And what they have noted was all the things which were there in the green represents the

responders.

That means, patient who has achieved the major pathological response and the red one represents

the patients who are non-responders.

That major pathological response was not achieved.

So, if you see this graph, in patient who has achieved major pathological response, the quantity or the ficile bacterium prosonage is enriched in the feces of these responders

when compared to the non-responders.

Similarly, if you see this graph, quantification of this ficile bacterium, it is significantly

higher in the patient who got the major pathological response when compared to the patient who

has not got the major pathological response.

So, in simple terms, patient who are responding to the immunotherapy has got high amount of

this ficile bacterium and patients who are not responding to the immunotherapy has got

less amount of this ficile bacterium.

And similarly, they have that isolated usage of this ficile bacterium alone does not had

slowed down the tumor growth.

But when it has been combined with the PD-1 monoclonal antibody, it enhances the anti-tumor

effect of the PD-L1 monoclonal antibody.

And the basic underlying mechanism was as I described previously, mainly because of the

valoric acid, which is a metabolite of this ficile bacterium, which will inhibit the interleukin

4 induced 1 gene, which in turn decreases the catabolism of tryptophan, thereby decreasing

the availability of the tryptophan metabolites, which in turn causing CD8 T cell activation,

which will help in the activation or potentially increasing the efficacy of immunotherapy.

So, again this is the key result, second result, what they have identified was the metabolite

valoric acid present in this ficile bacterium, it up regulates the run which in turn inhibits

the interleukin 4 induced gene 1.

And because of this interleukin 4 induced gene 1 is less, it inhibits the tryptophan

catabolism, thereby decreasing the tryptophan metabolite products and thereby again causing

increased activation of the CD8 T cell.

So, the conclusion of this study was yes, there is a potential for this ficile bacterium

in enhancing the PD-1 inhibitor efficacy.

And there are implications for personalised non-small cell lung cancer treatment strategies

with special emphasis on importance of identifying specific microbial metabolites. And thank you.